

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 194666

TO: Minh-Tam Davis Location: 3a24 / 3c18 Thursday, July 06, 2006

Art Unit: 1642

Phone: 571-272-0830

Serial Number: 09 / 403440

From: Jan Delaval

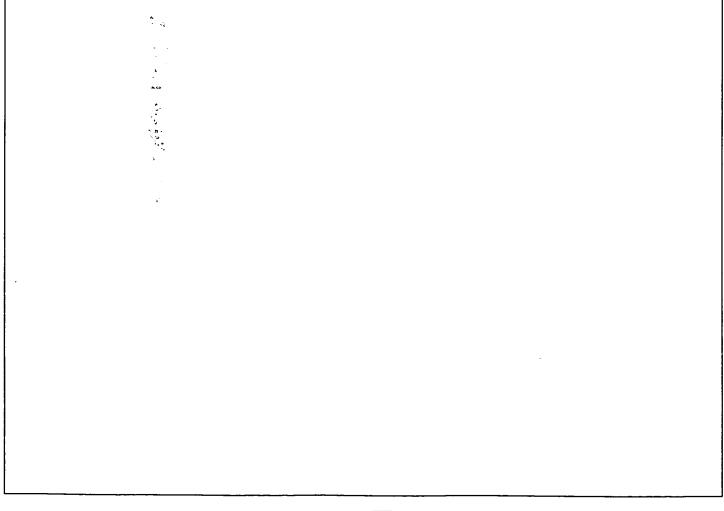
Location: Biotech-Chem Library

Remsen 1a51

Phone: 571-272-2504

jan.delaval@uspto.gov

Search Notes





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From:

Chan, Christina

Sent:

Wednesday, July 05, 2006 9:56 AM

To: Subject:

Davis, Minh-Tam; STIC-Biotech/ChemLib RE: Rush search request for 09/403440

Pleasemush Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 (571)-272-0841 Remsen, 3E89

-----Original Message-----

From:

Davis, Minh-Tam

Sent:

Monday, July 03, 2006 11:37 AM

To:

Chan, Christina

Subject:

Rush search request for 09/403440

Please search in commercial database, issued patent files and PGUB:

The peptide SEQ ID NO:2, with and without size limitation for the sequences in the database to the size of SEQ ID

NO:2.

Thank you.

MINH TAM DAVIS

ART UNIT 1642, ROOM 3A24, MB 3C18

272-0830

Searcher:__ Searcher Phone:_ Date Searcher Picked up:_ Date completed: 716106 Searcher Prep Time:_ Online Time:_

Т	ype of Search
NA#	AA#:_ <u>\</u>
	Oligomer:
Encode/	Transl:
Structure	e #:Text:
Inventor	· Litigation:

Vendors and cost where applicable	
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Result
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Maximum Match 100%
Listing first 45 summaries
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                 -hydroxybenzoate
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A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-393 <LAM>
A;Conserveferences: UNIPROT:P04637; UNIPARC:UPI000006F8F0; EMBL:X01405; GB:M13121; GB:N1
A;Buchman, V.L.; Chumakov, P.M.; Ninkina, N.N.; Samarina, O.P.; Georgiev, G.P.

R;Buchman, V.L.; Chumakov, P.M.; Ninkina, N.N.; Samarina, O.P.; Georgiev, G.P.

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R;Buchman, V.L.; Chumakov, P.M.; Samarina, N.N.; Samarina, N.N.

A; Molecule type: DNA A; Residues: 1-393 < BUC1>

A;Cross-references: UNIPARC:UPI00006F8F0; EMBL:M22898; NID:g189474

R;Lamb, P.; Crawford, L.

Mol. Cell. Biol. 6, 1379-1385, 1986

A;Title: Characterization of the human p53 gene.

A;Reference number: A25224; MUID:87064416; PMID:2946935

A; Accession: A25224

C;Species: Homo sapiens (man) C;Date: 05-Oct-1988 #sequence revision 18-Nov-1994 #text_change 09-Jul-2004 C;Accession: A55224; A43073; JT0436; S40773; S42869; A22837; A55080; A25397; B25397; S4 4905; I58354; I78850; I52681; S60153

cellular tumor antigen p53 [validated] - human N;Alternate names: cellular phosphoprotein p53; oncoprotein p53; transformation suppres

Qy 1 PPLSQETFSDLWKLLPEN 18	Query Match 94.3%; Score 99; DB 2; Lengthest Local Similarity 100.0%; Pred. No. 3.2e-07; Matches 18; Conservative 0; Mismatches 0; I	C;Genetics: A;Gene: p53 C;Superfamily: cellular tumor antigen p53 C;Keywords: tumor	A;Molecule type: mRNA A;Residues: 1-391 <lea> A;Cross-references: UNIPROT:Q95330; UNIPARC:UPI0000131047;</lea>	A; Title: cDNA cloning and immunological characterization A; Reference number: JC6193; MUID: 97208869; PMID: 9055811 A; Accession: JC6193	RiLe Goas, F.; May, P.; Ronco, P.; de Fromentel, C.C. Gene 185. 169-173, 1997	<pre>cumor suppressor pb3 - rabbit C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: 11-Apr-1997 #sequence_revision 09-May-1997 #text. C:Accession: .TCf193</pre>	ALIGNMENTS	44 41. 44 41.	N	44 41.9 239 2	44 41.9 211 1 44 41.9 238 2	44 41.9 209 2	44 41.9 114 2	41.9 113	44.5 42.4 211 2	45 42.9 837 2	45 42.9 405 2	45 42.9 367
	Length 391; ; ; Indels 0; Gaps 0;		; EMBL:X90592; NID:g1532043;	of rabbit p53.		#text_change 09-Jul-2004		protein - protein -	ner protein - simi NEF protein - simi	protein -	nef protein - simi	NEF protein - simi		conserved nypotnet L3 protein - bovin	conserved hypothet			hypothetical prote

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A;Molecule type: mRNA
A;Residues: 1-71, 'p', 73-393 <ZAK>
A;Residues: 1-71, 'p', 73-393 <ZAK>
A;Cross-references: UNIPARC:UPI000002ED67; EMBL:X02469; EMBL:M60950;
R;Harlow, E; Williamson, N.M.; Ralston, R.; Helfman, D.M.; Adams, T.
Mol. Cell. Biol. 5, 1601-1610, 1985
A;Title: Molecular cloning and in vitro expression of a cDNA clone fo A;Reference number: A55060; MUID:85267676; PMID:3894933
A;Accession: A55060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Residues: 1-71,'P',73-78,'T',80-393 <HAR2>
A;Cross-references: UNIPARC:UDI000031A30; EMBL:M14695; NID:g339815; PIDN:AAA61212.1;
A;Experimental source: clone p53-H-19, transformed hybridoma SV-80 cell line
R;Matlashewski, G.J.; Tuck, S.; Pim, D.; Lamb, P.; Schneider, J.; Crawford, L.V.
Mol. Cell. Biol. 7, 961-963, 1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: mRNA
A; Residues: 1-71, 'P', 73-272, 'H', 274-393 <HAR>
A; Residues: 1-71, 'P', 73-272, 'H', 274-393 <HAR>
A; Cross-references: UNIDARC: UPI000014ABB2; GB: K03199; NID: g189478;
A; Experimental source: clone pR4-2, cell line A431
R; Harris, N.; Brill, E.; Shohat, O.; Prokocimer, M.; Wolf, D.; Arai Mol. Cell. Biol. 6, 4650-4656, 1986
Mol. Cell. Biol. 6, 4650-4656, 1986
A; Title: Molecular basis for heterogeneity of the human p53 protein A; Reference number: A93086; MUID: 87089826; PMID: 3025664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Noce: this 72-Arg allele appears to be about 5 times more frequent than the A;Accession: JT0436
A;Molecule type: DNA
A;Residues: 1-71,'P',73-393 <BUC2>
                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPARC:UPI00001739B5
A;Experimental source: clone lambda C113
A;Note: 72-Cys was also found, and appear:
A;Accession: S42453
                                                                                                                                                                                                                                                                  A; Molecule type: mRNA; DNA
A; Residues: 66-79 < MKI3>
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EMBO J. 4, 1251-1255, 1985
A;Title: Human p53 cellular tumor antigen: cDNA sequence and expression A;Reference number: A22837; MUID:85230577; PMID:4006916
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A;Cross-references: UNIPARC:UPI000016AB44; EMBL:X01405; NID:g35215;
A;Zakut-Houri, R.; Bienz-Tadmor, B.; Givol, D.; Oren, M.
   A;Status:
                           A;Reference number: 138082; MUID:92007731;
A;Accession: 138082
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EMBO J. 3, 3257-3262,
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;Residues: 1-78,'T',80-393 <HAR1>;Cross-references: UNIPARC:UPI000014409D; EMBL:M14694; NID:g339813; PIDN:AAA61211.1;
;Experimental source: clone p53-H-1, transformed hybridoma SV-80 cell line
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                                                                                                Title: p53 is frequently mutated in Burkitt's lymphoma cell lines
                                                                                                                            Experimental source: clone J6K Farrell, P.J.; Allan, G.J.; Shanahan, IBO J. 10, 2879-2887, 1991
                                                                                                                                                                                                                                Cross-references: UNIPARC:UPI00001739B6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecule type: mRNA; DNA
Residues: 66-71,'P',73-79 <MKI2>
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translated from GB/EMBL/DDBJ
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A; MOLECULE type: mRNA
A; MOLECULE: 1-189, 'LISILSEWKEICVWSIWMTETLFDIVWWCPMSRLRLALT', 'VPPSTTTTCVTVPAWAA' <F01>
A; Cross-references: UNIPARC:UPI000011F824; EMBL:X60010; NID:9506432; PIDN:CAA42625.1;
A; Note: deletion of a C nucleotide causes a frameshift at position 566
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A;Accession: I38083 A;Status: translated from GB/EMBL/DDBJ

A; Residues: 1-192, 'R', 194-393 < F02>
A; Cross-references: UNIPARC: UPI0000070FCE; A; Molecule type: mRNA A; Accession: I38084 EMBL:X60011; NID:g506434; PIDN:CAA42626.1;

A,Status: translated from GB/EMBL/DDBJ A,Molecule type: mRNA A,Residues: 1-393 <F03>

A; Cross-references: UNIPARC: UPI000006F8F0; EMBL: X60012; NID:g506436; PIDN: CAA42627.1;

A;Accession: I38085 A;Status: translated from GB/EMBL/DDBJ

A; Molecule type: mRNA A; Residues: 1-245, T', 247-393 < F04> EMBL:X60013; NID:g506438; PIDN: CAA42628 .1;

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A;Cross-references: UNIPARC:UPI000006CFA4; A;Accession: I38086 A;Status: translated from GB/EMBL/DDBJ A;Molecule type: mRNA A;Residues: 1-236;'I',238-393 <F05> EMBL:X60014; NID:g506440;

A,Status: translated from GB/EMBL/DDBJ A,Molecule type: mRNA A,Residues: 1-247,'Q',249-393 <F06> A;Cross-references: UNIPARC:UPI0000072EB7; A;Accession: I38087 PIDN: CAA42629.1; ΡI

A;Cross-references: UNIPARC:UPI0000072FD4; A;Accession: I38088 EMBL: X60015; NID: g506442;

PIDN: CAA42630.1;

PI

A;Molecule type: mRNA
A;Residues: 1-71,'P',73-237,'Y',239-393 <FI
A;Cross-references: UNIPARC:UPI000006EDFC; A;Status: translated from GB/EMBL/DDBJ <F07> EMBL:X60016; NID:g506444; PIDN:CAA42631.1;

A;Status: translated from GB/EMBL/DDBJ A;Molecule type: mRNA A;Residues: 1-247,'Q',249-393 <F08> A;Cross-references: UNIPARC:UPI0000072FD4; A;Accession: I38090 A;Accession: I38089 A;Status: translate

A;Status: translated from GB/EMBL/DDBJ EMBL:X60017; NID:g506446; PIDN: CAA42632.1;

A;Molecule type: mRNA A;Residues: 1-71,'P',73-162,'H',164-393 <F09> A;Cross-references: UNIPARC:UPI000007055C; EM A; Accession: I38091 EMBL:X60018; NID:g506448; PIDN:CAA42633.1;

A;Status: translated from GB/EMBL/DDBJ A;Molecule type: mRNA A;Residues: 1-212,'Q',214-393 <F10> A; Cross-references: UNIPARC: UPI00000701E7; EMBL:X60019; NID:g506450; PIDN: CAA42634.1;

Status: translated from GB/EMBL/DDBJ

A;Status: Lienz: mRNA A;Molecule type: mRNA Cross references: UNIPARC:UPI000006E77E; EMBL:X60020; NID:g506452; PIDN:CAA42635.1; Note: all sequences submitted to the EMBL/GenBank/DDBJ databases June 1991

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R;Putreal, P.A.; Barrett, J.C.; Wiseman, R.W. Nucleic Acids Res. 19, 6977, 1991 A;Title: An Alu polymorphism intragenic to the TP53 gene A;Reference number: I38093; MUID:92107726; PMID:1762941

A;Residues: 1-393 <FUT> A; Molecule type: DNA A;Status: translated from GB/EMBL/DDBJ A; Accession: I38093

A;Cross-references: UNIPARC:UPI000006F8F0; EMBL:X54156; NID:g35213; PIDN:CAA38095.1; R;Yamada, Y.; Yoshida, T.; Hayashi, K.; Sekiya, T.; Yokota, J.; Hirohashi, S.; Nakata Cancer Res. 51, 5800-5805, 1991 A; Reference number: A44905; A; Accession: A44905 A; Title: p53 gene mutations in gastric cancer MUID:92034678; PMI PMID:1933850 in Nakatanı PII

A;ACCEBSTON.
A;ANOLECULE type: DNA
A;Residues: 246-247, W',249-250 <YAM>
A;Residues: 246-247, W',249-250 <YAM>
A;Cross-references: UNIPARC:UPI000011E80C; GB:S63157; NID:g237829; PIDN:AAB20140.1;
A;Cross-references: UNIPARC:UPI0000011E80C; GB:S63157; NCBIP:63158) PID

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Nucleic Acids Res. 17, 8375, 1989
A;Title: Nucleotide sequence of a cDNA encoding the monkey cellular phosphoprotein p53. A;Accession: S06594
A;Accession: S06594
A;Ccoss-references: UNIPROT: P13481; UNIPARC: UPTTC: C;Superfamily: cellular tumor antimor C;Keywords: apoptosis; Cellular tumor antimor F;392/m² (238.7%)
    RESULT
S51648
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A;Rosidues: 327-331, 'QCTSFQKENC' <CHO>
A;Residues: 327-331, 'QCTSFQKENC' <CHO>
A;Residues: 327-331, 'QCTSFQKENC' <GHO
A;Rosidues: mutant sequence with altered splicing and termination expressed in Molt-4 T-lymm R;Petersen, G; Song, D,; Huegle-Doerr, B.; Oldenburg, I.; Bautz, E.K.F.
Mol. Gen. Genet. 249, 425-431, 19-05
Mol. Gen. Genet. 249, 425-431, 19-05
A;Title: Mapping of linear epitopes recognized by monoclonal antibodies with gene-fragme A;Reference number: S60151; MUID:96133682; PMID:8552047
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A;Molecule type: DNA
A;Residues: 274-277,'S',279-282 <HEN2>
A;Cross-references: UNIPARC:UPI000011F7CD; GB:S41977; NID:g1679932; PIDN:AAB19325.1;
A;Crosw, V.T.; Quek, H.H.; Tock, E.P.C.
Cancer Lett. 73, 141-148, 1993
A;Title: Alternative splicing of the p53 tumor suppressor gene in the Molt-4 T-lympho
A;Reference number: I52681; WUID:94036762; PMID:8221626
A;Accession: I52681
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A;Title: Use of the single strand conformation polymorphism technique and A;Reference number: 158354; MUID:91296386; PMID:1648702
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A; Residues: 244-247, 'W', 249-252 <HENI>
A; Cross-references: UNIPARC: UPI000011F7CC; GB:S41969;
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                                                                                                                                                                                                                                                                             Superfamily: cellular tumor antigen p53;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; phosph;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; phosph;176,179,238,242/Binding site: zinc (Cys, His, Cys, Cys) #status predicted;392/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                 Cross-references: UNIPROT:P13481; UNIPARC:UPI0000131039; EMBL:X16384; NID:g22795; PID:
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; Pred. No. 3.2e-(
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                                                                C;Accession: JH0633
R;Legros, Y:, McIntyre, P.; Sc
Gene 112, 247-250, 1992
A;Title: The CDNA cloning and
A;Reference number: JH0633; Mt
                                       A; Reference number: A; Accession: JH0633
                                                                                                                                                                                 cellular tumor antigen p53 - golden hamster N;Alternate names: tumor-suppressor protein p53 C;Species: Mesocricetus auratus (golden hamster) C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999
A; Molecule type: mRNA
A; Residues: 1~396 < LE
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A;Molecule type: mRNA
A;Residues: 1-393 <LEES
A;Cross-references: UNIPROT:O09185; UNIPARC:UPI0000170734; GB:U50395; NID:g1842229; PID:
C;Comment: This protein is a multimer, it plays the central role in a complex DNA damaguiption, and recombination by protein/protein interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Lee, H.; Larner, J.M.; Hamlin, Gene 184, 177-183, 1997
A;Title: Cloning and characterizations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPROT:Q29628; UNIPARC:UPI0000131035; EMBL:X81704; NID C;Superfamily: cellular tumor antigen p53 C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; pho F;168,171,231,235/Binding site: zinc (Cys, His, Cys, Cys) #status predicted F;385/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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R;Dequiedt, F.; Willems, L.; Burny, A.; Kettmann, submitted to the EMBL Data Library, September 1994
A;Description: Nucleotide sequence of the ovine p5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Reference number: JC6176;
A;Contents: liver
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A;Molecule type: mRNA
A;Residues: 1-386 <DEQ>
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A; Accession: S51648
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;Alternate names: tumor-suppressor protein p53 ;Species: Bos primigenius taurus (cattle) ;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 10-Sep-1999 #s
                                                                                                                                                                                                                                                                                                                                                                                           Superfamily: cellular tumor antigen p53
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13 PLSQETFSDLWKLLPPN
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17; Conserv
                                                                                                                                                                                                                    Similarity
                                                                                   PLSOETFSDLWKLLPEN 18
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tulus griseus (Chinese hamster)
997 #sequence_revision 09-May-1997
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Pred. No.
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Pred. No. 1.8e-06;
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                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                              Length 393;
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P.; Soussi,

nd immunological MUID: 92210007;

PMID:1555773

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hamster

p53

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09-Jul-2004

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A;Gene: p53
C;Superfamily: cellular tumor antigen p53
C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; F;150,153,213,217/Binding site: zinc (Cys, His, Cys, Cys) #status predicted F;362/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 0 C;Accession: A29376; S61531; S72313; I51639 R;Soussi, T.; de Fromentel, C.C.; Mechali, M.; May, P.; Kress, M. Oncogene 1, 71-78, 1987 A;Title: Cloning and characterization of a cDNA from Xenopus laev A;Reference number: A29376; MUID:88143684; PMID:2830576
                                                                                                                                                                                                      RESULT
S38824
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A;Molecule type: mRNA
A;Residues: 1-51,'S',53-70,72-293,295-363 .
A;Cross references: UNIPARC:UPI0000171595;
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPARC:UPI0000171479; EMBL:X77546; NID:g468513; PIDN:CAA54672.1; R;Hoever, M.; Clement, J.; Wedlich, D.; Montenarh, M.; Knochel, W. submitted to the EMBL Data Library, March 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: Overexpression of wild-type p53 interferes with normal development A;Reference number: I51639; MUID:94134403; PMID:8302570 A;Accession: S61531
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A; Accession: S72313
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Oncogene 9, 109-120, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-363 < SOU>
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A; Accession: A29376
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C;Species: Xenopus laevis (African clawed frog)
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                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
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82.4%;
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94.1%;
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Pred. No. 0
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; EMBL:X77546; NID:g468513; PIDN:CAA54672.1;
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                              alternative splicing
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                                                                                                                                 23-Jul-1999
                                                                             O.; Rotter,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 N;Alternate names: Oncoprotein p53
C;Species: Mus musculus (house mouse)
C;Date: 28-Aug-1985 #sequence revision 04-Oct-1996 #text change 09-C;Date: 28-Aug-1985 #sequence revision 04-Oct-1996 #text change 09-C;Accession: A22739; S06336; AD2684; S38822; S38823; S40014; I48703 R;Blenz, B; Zakut-Houri, R.; Givol, D.; Oren, M.
BMBO J. 3, 2179-2183, 1984
A;Title: Analysis of the gene coding for the murine cellular tumour A;Reference number: A22739; MUID:85027173; PMID:6092064
A;Accession: A22739
                         A;Cross references: UNIPARC:UDIO001449CC
R;Zakut-Houri, R.; Oren, M.; Bienz, B.; Lav
Nature 306, 594-597, 1983
A;Title: A single gene and a pseudogene for
A;Reference number: A02684; MUID:84068204;
A;Accession: A02684
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C;Comment: This sequence, produc
s not know.
C;Superfamily: cellular tumor an
C;Keywords: alternative splicing
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A;Title: Alternatively spliced p53 RNA in transformed and
A;Reference number: S35478; MUID:92253421; PMID:1579500
                                                                                                                                                                                                                                                     Bioorg. Khim. 13, 1691-1694, 1987
A;Title: Primary structure of DNA complementary to murin
A;Reference number: S06336; MUID:88221682; PMID:3329909
A;Accession: S06336
                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-134, 'V', 136-390 <BIE>
A; Cross-references: UNIPROT: P02340;
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A;Residues: 1-381 <HAN>
A;Cross-references: UNIPARC:UPI000016CF91;
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A;Molecule type: mRNA
A;Residues: 1-381 <ARA>
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F;77,9,12,18,23,37/Binding site: phosphate (Ser) (covalent) #status predicted
F;173,176,235,239/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;312/Binding site: phosphate (Ser) (covalent) (by cdc2 kinase) #status predi
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                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-134,'V',136-390 < CHU>
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Best Local S
Matches 14
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       type: mRNA
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     016CF91; EMBL:M13874; NID:g200202; PIDN:AAA39883.1; s submitted to the EMBL Data Library, July 1988 by alternative splicing of the tenth intron, lacks
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                                                                                                                                                                                                                                   translation
                                                        for the cellular
)4; PMID:6646235
                                                                                                                                 Lavie, V.; Hazum,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2;
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                                                                                                                                                                                                                                                                                                              murine
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                                                                                tumour antigen p53
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N;Alternate names: gene p53 protein; nuclear oncoprotein C;Species: Rattus norvegicus (Norway rat) C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_C;Accession: S02192; S41149 C;Accession: S02192; S41149 C;C; Breugnot, C.; May, E. R;Soussi, T.; de Fromentel, C.C.; Breugnot, C.; May, E. Nucleic Acids Res. 16, 11384, 1988 A;Title: Nucleotide sequence of a cDNA encoding the rat pA;Reference number: S02192; MUID:89083585; PMID:3060862 A;Accession: S02192
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A;Residues: 1-167,'G', 169-390 <ARA3>
A;Cross-references: UNIPARC:UPI000016CF90; EMBL:M13873;
A;Cross-references: UNIPARC:UPI000016CF90; EMBL:M13873;
A;Jenkins, J.R.; Rudge, K.; Redmond, S.; Wade-Evans, A.
Nucleic Acids Res. 12, 5609-5626, 1984
A;Title: Cloning and expression analysis of full length
A;Reference number: 148703; MUID:84272240; PMID:6379601
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A;Residues: 1-167, 'G',169-233,'I',235-390 <ARA2>
A;Residues: 1-167, 'G',169-233,'I',235-390 <ARA2>
A;Cross-references: UNIPARC: UPI00001739B9; EMBL.M13873
R;Arai, N.; Nomura, D.; Yokota, K.; Wolf, D.; Brill, E.; Shohat, submitted to the EMBL Data Library, July 1988
A;Reference number: S40014
A;Accession: S40014
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cellular tumor antigen p53 - rat
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F;168-178/Region: conserved region III
F;231-252/Region: conserved region IV
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A;Cross-references: UNIPARC:UPI00001739B8; GB:X01237; GB:K01700; NID:g53575
R;Arai, N., Nomura, D., Yokota, K.; Wolf, D.; Brill, E.; Shohat, O.; Rotter, Mol. Cell. Biol. 6, 3232-3239, 1986
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Best Local
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114-139/Region: conserved region II
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;Residues: 1-390 <ARAl>
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Pred. No. 0.0019;
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glutamate receptor delta-1 chain precursor - rat N,Alternate names: kainate receptor C;Species: Rattus norvegicus (Norway rat) C;Date: 07-Apr-1994 #sequence_revision 07-Apr-199 C;Accession: $28857; 331222 R;Lomeli, H.; Sprengel, R.; Laurie, D.J.; Koehr, FEBS Lett. 315, 318-322, 1993 A;Title: The rat delta-1 and delta-2 subunits ext
                                 A;Recession: S28857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Comment: This protein is the product of a tumor suppressor gene, p53, who: C;Superfamily: cellular tumor antigen p53
C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nuce;164,167,227,231/Binding site: zinc (Cys, His, Cys, Cys) #status predicted F;395/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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A;Residues: 1-396 <DEF>
A;Residues: 1-396 <DEF>
A;Cross-references: UNIPROT:P25035; UNIPARC:UPI0000131043; GB:M75145; NID:g213828; PIDN
A;Experimental source: liver
A;Experimental source: liver
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C;Species: Oncorhynchus mykiss (rainbow trout)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 09-Jul-2004
C;Accession: JH0631
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Nucleic Acids Res. 21, 713-717, 1993

A;Title: Structure of the rat p53 tumor suppressor gene.

A;Reference number: S41149; MUID:93181268; PMID:8441680
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Gene 112, 241-2
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C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nu F;174,177,236,240/Binding site: zinc (Cys, His, Cys, Cys) #status predicted F;390/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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A;Residues: 1-173,'W',175-391 <HUL>
A;Cross-references: UNIPARC:UP100001679E9; EMBL:L07909
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A;Cross-references: UNIPROT:P10361; UNIPARC:UPI0000131048; EMBL:X13058; NID:g56828; PID
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Matches
Molecule type: mRNA
                     Status: nucleic acid sequence
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Pred. No. 0.0039;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 60;
Pred. No.
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precorrin-2 methyltransferase homolog - Methanococcus jannaschii C,Species: Methanococcus jannaschii C,Species: Methanococcus jannaschii G,Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 05-Oct-2004 C;Accession: C64396 R,Bult, C,J; Weinstock, K,G.: Merrick JM,; Reich, C.I.; Overbeek, R.; Kirkness, E.F.; Weinstock, K,G.: Merrick JM,
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A;Residues: 1-1009 <XAM>
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A;Rolecule type: maintenance for the first status predicted cattle properties of transmembrane protein
A;Cross-references: UNIPARC:UPI0000170981; EMBL:Z17238; NID:g56285; PIDN:CAA78936.1; PID
C;Superfamily: mannose for the receptor cattle receptor; transmembrane protein
C;Keywords: glycoprotein; ion channel; neurotransmitter receptor; transmembrane protein
F;1-15/Domain: signal sequence #status predicted cattle predicted cattle
F;161-109/Product: glutamate receptor homology cattle
F;161-86/Domain: transmembrane #status predicted cattle
F;565-884/Domain: transmembrane #status predicted cattle
F;601-621/Domain: transmem
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A;Accession: S31222
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;422/Binding site: carbohydrate (Asn) (covalent) #status predicted
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               G.J.; Zhou, L.; Fleischmann, R.D.; Sutton, G.G.; Bla.
rkness, E.F.; Weinstock, K.G.; Merrick, J.M.; Glodek,
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Science 273, 1058-1073, 1996
A;Authors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.; Woese, C.A;Authors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.; Woese, C.A;Title: Complete genome sequence of the methanogenic archaeon, Methanococcus jannaschii A;Reference number: A64300; MUID:96337999; PMID:8688087
A;Accession: C64396
A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA
A;Residues: 1-230 <BUL-
A;Residues: 1-230 <BUL-
                                                                                                                                                                                                                                                                                                                          R;Bolotin, A.; Wincker, P.; Mauger, S.; Jaillon, Genome Res. 11, 731-753, 2001
A;Title: The complete genome sequence of the lac
                                                                                                                                                                                                                                                                                                                                                                                     C; Species: Lactococc
C; Date: 23-Mar-2001
C; Accession: G86715
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G86715
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A;Start codon: GTG
C;Superfamily: precorrin-2 C20-methyltransferase
                                Ś
                                                                                                                                                                                                                                                                                                                                                                                                     racemase [imported] - Lactococcus lactis subsp. lactis (strain IL1403)
C;Species: Lactococcus lactis subsp. lactis
C;Date: 23-Mar-2001 #sequence_revision 23-Mar-2001 #text_change 09-Jul-2004
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A; Residues: 1-367 <STO>
                                                                                                                                                                                                                                                                                                      A;Title: The complete genome sequence of the lactic acid bacterium Lactococcus A;Reference number: A86625; MUID:21235186; PMID:11337471
                                                                                                                                                      A;Gene: yhdB
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                                                                                                                                                                                                                                                                   A; Status: preliminary
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                                                                                                             Query Match
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GG; GC:00046872; F:metal ion binding; IE
GG; GC:0003700; F:transcription factor
GG; GC:0008270; F:zinc ion binding; IEA
GG; GC:0006915; P:apoptosis; IEA.
GG; GC:0006915; P:apoptosis; IEA.
GG; GC:0007049; P:cell cycle; IEA.
GG; GC:000535; P:negative regulation of
GG; GC:0006355; P:regulation of transcr
GG; GC:0006350; P:transcription; IEA.
InterPro; IPR011615; P53_DNA_bd.
Pfam; PF00870; P53; 1.
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PubMed=16131611; DOI=10.1101/gad.1339905;

Bourdon J.C., Fernandes K., Murray-Zmijewski F., Liu G., D
Xirodimas D.P., Saville M.K., Lane D.P.;

"p53 isoforms can regulate p53 transcriptional activity.";

Genes Dev. 19:2122-2137(2005).
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O; GO:0006915; P:apoptosis; IEA.
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                                                                                       UniProt Consortium,
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                                                                                                                                                                                            expression.

COPACTOR: Binds 1 zinc ion per subunit (By similarity).

SUBUNIT: Binds DNA as a homotetramer. Found in a complex with

CABLES1 and p53/TP73. Interacts with histone acetyltransferases

EP300 and methyltransferases HRWT1L2 and CARM1, and recruits them

to promoters. C-terminus interacts with TAP1, when TAP1 is part of

the TFIID complex. Interacts with HIPK1, HIPK2, AXIN1, and

P53DINP1. Part of a complex consistiting of TP53, HIPK2 and AXIN1.

Interacts with WMOX (By similarity).

SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity).

SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity).
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GO:0006915; P:apoptosis; IEA.
GO:0007049; P:cell cycle; IEA.
GO:0045786; P:negative regulation of progression through . . .
GO:0006355; P:regulation of transcription, DNA-dependent; IEA.
GO:0006350; P:transcription; IEA.
                                                     SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity). Prim: Phosphorylated. Phosphorylation on Ser residues mediates transcriptional activation. Phosphorylated on Thr-18 by VRK1, which may prevent the interaction with MDM2. Phosphorylated by HIPKI (By similarity). Prim: Acetylated. Its deacetylation by SIRT1 impairs its ability.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases appoptosis induction seems to be mediated either by simulation of appoptosis induction seems to be mediated either by simulation of appoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BAX and FAS antigen expression,
induce proapoptotic program
similarity)
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100.0%; Pr
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GO; GO:0005739; C:mitochondrion; ISS.
GO; GO:0005739; C:mitochondrion; ISS.
GO; GO:0005730; C:mutoledum; ISS.
GO; GO:0005524; F:ATP binding; ISS.
GO; GO:0005507; F:COpper ion binding; ISS.
GO; GO:0005507; F:CNA binding; ISS.
GO; GO:0005515; F:DNA binding; ISS.
GO; GO:0006215; P:apoptosis; ISS.
GO; GO:0006215; P:apoptosis; ISS.
GO; GO:0006215; P:apoptosis; ISS.
GO; GO:0006215; P:apoptosis; ISS.
GO; GO:0006215; P:caspase activation via
GO; GO:0007569; P:cell aging; ISS.
GO; GO:00007509; P:cell aging; ISS.
GO; GO:00007509; P:cell aging; ISS.
GO; GO:0000154; P:cell aging; ISS.
GO; GO:000154; P:cell aging; ISS.
GO; GO:000155; P:cell aging; ISS.
GO; GO:00015; P:cell aging; ISS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SMR; Q9TUB2; 86-282.
30; G0:0005739; C:mucleolus; ISS.
30; G0:0005730; C:mucleolus; ISS.
30; G0:0005730; C:mucleolus; ISS.
30; G0:0005524; F:ATP binding; ISS.
30; G0:0005527; F:copper ion binding; ISS.
30; G0:0003677; F:DNA binding; ISS.
30; G0:0006739; F:DNA strand annealing activity;
30; G0:0006739; F:protein binding; ISS.
30; G0:0006915; P:apoptosis; ISS.
30; G0:0006915; P:apoptosis; ISS.
30; G0:0006915; P:cappasse activation via cytochr.
30; G0:0006759; P:cell aging; ISS.
30; G0:0007569; P:cell cycle arrest; ISS.
30; G0:0007059; P:cell proliferantion; ISS.
30; G0:0007054; P:cell proliferantion; ISS.
30; G0:00030154; P:cell proliferantion; ISS.
30; G0:00030154; P:cell proliferantion; ISS.
30; G0:00030154; P:cell proliferantion; ISS.
30; G0:0000283; P:cell proliferantion of cell gr
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SIMILARITY: Belongs to the p53 family.
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transformed cells.
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llarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P:apoptosis; ISS.
P:base-excision repair; ISS.
P:caspase activation via cytochrome
P:cell aging; ISS.
P:cell cycle arrest; ISS.
P:cell differentiation; ISS.
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                                                                                                                                              MW.
                                                                                                                                                                                                                Zinc (By similarity).
S'-phospho-RNA (covalent) (()
similarity).
Phosphoserine (by PRPK) (By
Phosphothreonine (by VRKI)
similarity).
29
                                                                      ۰,
                                                                                       Score 99;
Pred. No
                                                                                                                                                             N6-acetyllysine
N6-acetyllysine
N6-acetyllysine
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Cellular tumor antigen
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Nuclear export signal
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Interaction with HIPK1
Interaction with AXIN1
Interaction with HIPK2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /FTId=PRO_0000185710
By similarity
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                                                                                                                                      6-acetyllysine (By similarity)
6-acetyllysine (By similarity)
6-acetyllysine (By similarity)
A4C3D88E8DF55162 CRC64;
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Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      activity;
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localization signal
                                                                                                       Length 386;
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P53 DELLE STANDARD; P; Q885Z3; 20-DEC-2005, integrated into Un 01-JUN-2002, sequence version 1 07-FEB-2006, entry version 22. Cellular tumor antigen p53 (Tum
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MEDLINE=22030464; PubMed=12034505; DOI=10.1016/S0378-1119(02)00472-9;

Xu N., Shiraki T., Yamada T., Nakajima M., Gauthier J.M.,

Pfeiffer C.J., Sato S.;
                                                                                                                                                                                                                                                                        Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                       Copyrighted
GO: GO:0005739; C:mitochondrion; ISS.
GO: GO:0005730; C:nucleolus; ISS.
GO: GO:0005524; F:AIP binding; ISS.
GO:0005524; F:Copper ion binding; ISS.
GO:0005507; F:DNA binding; ISS.
GO:0003677; F:DNA binding; ISS.
GO:00005515; F:Drotein binding; ISS.
GO:0005515; F:protein binding; ISS.
GO:0006915; P:apoptosis; ISS.
GO:0006814; P:base-excision repair; ISS.
GO:0006284; P:base-excision via cytochrome; GO:0007569; P:cell aging; ISS.
TO: GO:0007569; P:cell cycle arrest; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE [MRNA]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=9749;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Odontoceti; Monodontidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Cetacea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Delphinapterus leucas (Beluga whale)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression.

expression.

COFACTOR: Binds 1 zinc ion per subunit (by similarity).

COFACTOR: Binds DNA as a homotetramer. Found in a complex with SUBUNIT: Binds DNA as a homotetramer. Found in a certyltransferases EP300 and methyltransferases HRMTIL2 and CARMI, and recruits the to promoters. C-terminus interacts with TAFI, when TAFI is part the TFIID complex. Interacts with HIFK1, HIFK2, AXINI, and P53DINP1. Part of a complex consistiing of TP53, HIFK2 and AXIN Interacts with WMOX (By similarity).

Thereacts with WMOX (By similarity).

Thereacts with MWOX (By similarity).

Thereacts with WMOX (By similarity).
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                                                                                                                                                                                                                                                                                                                similarity). is found in increased amounts in transformed cells. p53 is frequently mutated c many types of cancer. SIMILARITY: Belongs to the p53 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of
                                                                                                                                                                                                                                                                                                                                                                                                                                 PTM: Phosphorylated. Phosphorylation on Ser residues mediates transcriptional activation. Phosphorylated on Thr-18 by VRKI, which may prevent the interaction with MDM2. Phosphorylated by HIPKI (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                          induce
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Q9NP68; 1AIE.
Q8SPZ3; 87-283.
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Matches 18
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                                                                                                                                                                                                                 Q9W\u00fcR6;
Q9W\u00fcR6;
Q1-DEC-2000, integrated into U
01-NCV-1999, sequence version 46.
07-FEB-2006, entry version 46.
(The control of the control of the
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MOD_RES
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METAL
TISSUE=Spleen;
MEDLINE=99265972; PubMed=10331945; DOI=10.1006/geno.1999.5794;
n'erchia A.M., Pesole G., Tullo A., Saccone C., Sbisa E.;
                                                                                                                   Eukaryota; Metazoa; Chordata; Craniat; Mammalia; Eutheria; Euarchontoglires; Hystricognathi; Caviidae; Cavia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGION
REGION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGION
                                                                                                                                                     Cavia porcellus (Guinea pig).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                         P53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Acetylation; Activator; Anti-oncogene; Apoptosis; Cell cycle; DNA-binding; Metal-binding; Nuclear protein; Phosphorylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ProDom; PD002681; P53; 1. PROSITE; PS00348; P53; 1.
                                                                 NUCLEOTIDE SEQUENCE [MRNA]
                                                                                                 NCBI_TaxID=10141;
                                                                                                                                                                                               Name=TP53;
                                                                                                                                                                                                             Cellular tumor antigen p53
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172
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109
313
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362
                                                                                                                                                                                                                                                                    integrated into UniProtKB/Swiss-Prot.
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1 387 Cellular tumor antigen
/FTId=PRO_000185699.
                                                                                                                                                                                                                                                                                                       STANDARD;
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P:negative regulation of cell growth; ISS
P:nucleotide-excision repair; ISS.
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N6-acetyllysine (By similarity).
N6-acetyllysine (By similarity).
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5'-phospho-RNA (covalent) (By
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N6-acetyllysine
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Basic (repression of DNA-binding) (By
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                                                                                                                                          Rodentia;
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K1) (By
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Pfam; PF00
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SMR;
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         DNA_BIND
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                                                                                                                                                                                                                                    InterPro;
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               100
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"Guinea pig p53 mRNA: identification of new elements in coding and untranslated regions and their functional and evolutionary implications.";
Genomics 58:50-64(1999).
                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00870; P53; 1.
Pfam; PF07710; P53 tetramer; 1.
PRINTS; PR00386; P53SUPPRESSR.
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                                                                                      Acetylation; Activator; Anti-oncogene; Apoptosis; Cell cycle; DNA-binding; Metal-binding; Nuclear protein; Phosphorylation; Transcription; Transcription regulation; Zinc.
CHAIN 1 391 Cellular tumor antigen p53.
                                                                                                                                                                                                                                                                                                          ProDom; PD002681; P53; 1. PROSITE; PS00348; P53; 1.
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-!- SIMILARITY: Belongs to the p53 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30; GO:0005739; C:mitochondrion; ISS.
30; GO:0005739; C:mucleolus; ISS.
30; GO:0005524; F:ATP binding; ISS.
30; GO:0005524; F:ATP binding; ISS.
30; GO:0005507; F:copper ion binding; ISS.
30; GO:0005515; F:DNA binding; ISS.
30; GO:0005515; F:protein binding; ISS.
30; GO:0005515; F:protein binding; ISS.
30; GO:0006815; P:apoptosis; ISS.
30; GO:0006828; P:apoptosis; ISS.
30; GO:0006828; P:caspase activation via cyt
30; GO:000759; P:cell aging; ISS.
30; GO:000759; P:cell differentiation; ISS.
30; GO:0007050; P:cell differentiation; ISS.
30; GO:0030154; P:cell proliferation; ISS.
30; GO:00301828; P:negative regulation of cell
30; GO:0006289; P:negative regulation repair
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FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of max antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP300 and methyltransferases HRMT1L2 and CARM1, and recruits them to promoters. C-terminus interacts with ThF1, when ThF1 is part of the TFIID complex. Interacts with HIPK1, HIPK2, AXIN1, and P53DINP1. Part of a complex consisiting of TP53, HIPK2 and AXIN1. Interacts with WWOX (By similarity).

SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity). SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity). Fine Phosphorylated. Phosphorylated on Thr-18 by VRK1, which may prevent the interaction with MDM2. Phosphorylated by HIPK1. Phosphorylated on Ser-46 by HIPK2 upon UV irradiation. Phosphorylated on Ser-46 is required for acetylation by CREBBP (Proceedings).
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DISEASE: p53 is found in increased
Frankformed cells. p53 is frequent
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COFACTOR: Binds 1 zinc ion per subunit (By similarity).

SUBUNIT: Binds DNA as a homotetramer. Found in a complex with SUBUNIT: Binds DNA as a homotetramer. Found in a complex with CABLES1 and p53/TP73. Interacts with histone acetyltransferases can be completed by the complex of the complex with the complex with the complex and recruits the complex can be completed by the completed by the completed by the complex can be completed by the compl
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IPR011615; P53 DNA bd.
IPR012346; P53 RUNT DNA bd.
IPR010991; p53 tetrameristn.
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P:negative regulation of cell
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/FTId=PRO 0000185696.
By similarity.
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P53 RABIT
Q95330;
01-NOV-1997,
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SEQUENCE
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07-FEB-2006, entry version 53.
Cellular tumor antigen p53 (Tumo
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"cDNA cloning and immunological
                                                                                                                                                                                                             MEDLINE=97208869;
                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE (MRNA).
STRAIN=New Zealand;
                                                                                                                                                                                                                                                              Oryctolagus
                                                                                                                                                                                                                                                                       Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                            Oryctolagus
                                                                                                                                                                                                                                                                                                       Name=TP53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOTIF
                                                                                                                                                                                                                                                                                                                                                                                                                 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                PPLSOETFSDLWKLLPEN
                                                                                                                                                                                                                                                                                                                                                                                                                          PPLSQETFSDLWKLLPEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          380
391
                                                                                                                                                                                                                                                                                   cuniculus (Rabbit).
Metazoa; Chordata; (
                                                                                                                                                                                                                                                                                                                                                                                                                                                     94.3%;
larity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                           integrated into UniProtKB/Swiss-Prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ŗ,
                                                                                                                                                                                                                                                                                                                                                                 STANDARD
                                                                                                                                                                                          PubMed=9055811; DOI=10.1016/S0378-1119(96)00604-X; , Ronco P., Caron de Fromentel C.; immunological characterization of rabbit p53.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46
303
371
380
43288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     108
368
290
290
358
358
319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   390
                                                                                                                                                                                                                                                                       Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Glires; Lagomorpha; Leporidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ¥.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Interaction with HIPK1 (Interaction with AXINI (Interaction with HIPK2 (Bollgomerization, Basic (repressional Repression HIPK2)
                                                                                                                                                                                                                                                                                                               (Tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         similarity).

Binilarity).

Phosphoserine (by HIPK2) (By simil N6-acetyllysine (By similarity).

N6-acetyllysine (By similarity).

N6-acetyllysine (By similarity).

N8-321D40702383573E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 99; DB; Pred. No. 2.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zinc (By similarity).
5'-phospho-RNA (covalent) (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                  18
                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bipartite no
similarity)
Nuclear exp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Phosphothreonine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Phosphoserine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Transcription activation (acidic)
                                                                                                                                                                                                                                                                                                              suppressor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  export
                                                                                                                                                                                                                                                                                                                                                               391 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.8e-06;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ξġ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  signal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of DNA-binding).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 391;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRPK) (By
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VRK1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ( (By similarity).
(1 (By similarity).
(1 (By similarity).
(2 (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (By
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  similarity)
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EMBL; X90592; CAA62216.1; -
PIR; JC6193; JC6193.
HSSP; P04637; IGZH.
SMR; Q95330; 91-286.
GO; GO:0005739; C:mucleolus
GO; GO:0005739; C:mucleolus
GO; GO:0005507; F:CORPETION
GO; GO:0003677; F:DNA bindi
GO; GO:0003677; F:DNA bindi
GO; GO:0003677; F:DNA bindi
GO; GO:0003675; F:DNA bindi
GO; GO:0003675; F:DNA bindi
GO; GO:0003675; F:DNA bindi
GO; GO:0006915; F:Drotein b
GO; GO:0006915; F:Dapoptosis
GO; GO:0006915; P:Caspase
GO; GO:0007628; P:Caspase
GO; GO:0007569; P:Cell agin
GO; GO:0007569; P:Cell pirol
GO; GO:0008283; P:Cell pirol
GO; GO:0008283; P:Cell pirol
GO; GO:0008283; P:Cell pirol
GO; GO:0008283; P:Cell pirol
GO; GO:0008289; P:mucleotid
MOD_RES
MOD_RES
MOD_RES
SEQUENCE
                                                                                                                                                                                                         DNA BIND
REGION
REGION
REGION
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REGION
REGION
                                                                                                                                           METAL
METAL
                                                                                                                                                                    MOTIF
                                                            MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00870; P53; 1.
Pfam; PF07710; P53 tetramer; 1.
PRINTS; PR00386; P53SUPPRESSR.
                                                                                                     BINDING
                                                                                                                                                                                              MOTIF
                                                                                                                                                                                                                                                                                                                                                   Acetylation; Activator; Anti-oncogene; Apoptosis; Cell cycle; DNA-binding; Metal-binding; Nuclear protein; Phosphorylation;
                                                                                                                                                                                                                                                                                                                                                                              ProDom; PD002681; P53; 1. PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR011615; P53_DNA_bd.
InterPro; IPR012346; P53_RUNT_DNA_bd.
InterPro; IPR010991; p53_tetramerIstn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR002117;
InterPro; IPR011615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted
Distributed
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                                                                                                                                                                                                                                                                                                                                        Franscription;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30; GO:0005739; C:mitochondrion; ISS.
30; GO:0005730; C:nucleolus; ISS.
30; GO:0005524; F:ATP binding; ISS.
30; GO:0005527; F:copper ion binding; ISS.
30; GO:0005507; F:DNA binding; ISS.
30; GO:000759; F:DNA binding; ISS.
30; GO:0007515; F:protein binding; ISS.
30; GO:0005515; F:protein binding; ISS.
30; GO:000544; P:base-excision repair; ISS.
30; GO:0006284; P:base-excision repair; ISS.
30; GO:0006284; P:base-excision repair; ISS.
30; GO:0006284; P:call aging; ISS.
30; GO:0007569; P:cell cycle arrest; ISS.
30; GO:0007569; P:cell differentiation; ISS.
30; GO:0007589; P:cell proliferation; ISS.
30; GO:00030154; P:cell proliferation; ISS.
30; GO:00030154; P:nucleotide-excision repair.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             many types of cancer.
SIMILARITY: Belongs to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PTM: Phosphorylated. Phosphorylation on Ser residues transcriptional activation. Phosphorylated on Thr-18 which may prevent the interaction with MDM2. Phospho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DISEASE: p53 is found transformed cells. p5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PTM: Acetylated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              induce proapoptotic program
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      which may prevent the interaction HIPK1 (By similarity)
                                                                                                                                                                                                         99
63
97
113
317
323
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      by the UniProt
                                                                                                                                                                                                                                                                                                                                       Transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P:cell cycle arrest; ISS.
P:cell differentiation; ISS.
P:cell proliferation; ISS.
P:negative regulation of cell
P:nucleotide-excision repair;
                                                                                                     348
173
176
235
239
390
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385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        p53
  WW.
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                                                                                                                                                                                                                                /FTId=PRO_0000185711.

By similarity.
Transcription activation (
Interaction with WWOX (By
Interaction with HIPKI (By
Interaction with AXIN1 (By
Interaction with HIPK2 (By
              similarity).
N6-acetyllysine
N6-acetyllysine
N6-acetyllysine
                                                              similarity).
Phosphoserine (b)
Phosphothreonine
                                                                                                    Zinc (By similarity).
Zinc (By similarity).
5'-phospho-RNA (covalent)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA.
                                                                                                                                                                                          Basic (repression Bipartite nuclear
                                                                                                                                                                                                                                                                                                                         regulation; Zinc.
Cellular tumor antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     increased
                                                                                                                                                                    Nuclear
                                                                                                                                                                                 similarity)
                                                                                                                                                                                                                      Oligomerization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        frequently
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              p53 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and
                                                                                                                                         (By similarity)
(By similarity)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               iss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modulate cell senescence (By
                                                                                                                                                                  export signal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        see http://www.uniprot.org/terms
Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amounts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mounts in mutated o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytochrome
                                                                             ξģ
                                                                                                                                                                                                                                 h WWOX (By similarity).
h HIPK1 (By similarity)
h AXIN1 (By similarity)
h HIPK2 (By similarity)
             (By
VB)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               growth;
                                                               (by VR
                                                                                                                                                                                           of DNA-binding).
localization signal
              similarity)
similarity)
                                        similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 residues mediates
n Thr-18 by VRK1,
Phosphorylated b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or
                                                                  VRK1)
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                                                                                                                                                                   βу
                                                                                                                                                                                                                                                                                                                         p53.
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                                                                             βу
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                                                                                                      (By
                                                                                                                                                                                                                                                                                      (acidic)
                                                                                                                                                                    similarity)
                                                                                                                                                                                                                                   / similarity)
/ similarity)
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                                                               y similarity).
(By
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Best Local S
Matches 18
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P53_C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P13481,
EMBL; X16384; CAA34420.1; -; mRNA. PIR; S06594; S06594. HSSP; P04637; 10LG. SMR; P13481; 94-289. GO; GO:0005739; C:mitochondrion; ISS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cercopithecus aethiops (Green monkey) (Grivet).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JAN-1990, sequence vers
07-FEB-2006, entry version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cercopithecidae; Cercopithecinae; Cercopithecus. NCBI_TaxID=9534;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Name=TP53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JAN-1990,
                                                                                                         Distributed under the
                                                                                                                         Copyrighted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Nucleotide sequence of a cDNA encoding the monkey cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rigaudy P., Eckhardt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=90045967; PubMed=2530498;
                                                                                                                                                      transformed cells. p53 is frequently m many types of cancer.
SIMILARITY: Belongs to the p53 family.
                                                                                                                                                                                                                                                                                                               SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity). PTM: Phosphorylated. Phosphorylation on Ser residues mediates transcriptional activation. Phosphorylated on Thr-18 by VRK1, which may prevent the interaction with MDM2. Phosphorylated on Thr-55 by TAF1 which promotes MDM2-mediated p53 degradation. Phosphorylated by HIPK1 upon Phosphorylated by HIPK1. Phosphorylated on Ser-46 by HIPK2 upon
                                                                                                                                                                                                                                                                                                                                                                                                                     CABLES1 and p53/TP73. Interacts with histone acetyltransferases EP300 and methyltransferases HRWT1L2 and CARMI, and recruits them to promoters. C-terminus interacts with TAF1, when TAF1 is part of the TFIID complex. Interacts with HIPK1, HIPK2, AXIN1, and p53DINP1. Part of a complex consisting of TP53, HIPK2 and AXIN1. Interacts with WMOX (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression.

COFACTOR: Binds 1 zinc ion per subunit (By similarity)

SUBUNIT: Binds DNA as a homotetramer. Found in a compl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 leic Acids Res. 17:8375-8375(1989).

FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CERAE
                                                                                                                                                                                                           DISEASE: p53 is found in
                                                                                                                                                                                                                                                                by CREBBP (By similarity).
PTM: Acetylated. Its deacetylation by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BAX and FAS antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                              induce proapoptotic
                                                                                                                                                                                                                                                                                                   irradiation. Phosphorylation on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              controlling a set of genes required for this process. One of activated genes is an inhibitor of cyclin-dependent kinases, optosis induction seems to be mediated either by stimulation or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE [MRNA]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                           by the UniProt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         integrated into UniProtKB/Swiss-Prot.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94.3%;
100.0%;
                                                                                                         Creative Commons
                                                                                                                                                                                                                                                program
                                                                                                                         Consortium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 99;
Pred. No.
                                                                                                                                                                                                           increased amounts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                                and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                modulate cell
                                                                                                                                                                                                                                                                                                   Ser-46 is required
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            393
                                                                                                         see http://www.uniprot.org/terms
Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; L
2.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A
                                                                                                                                                                                             mutated
                                                                                                                                                                                                                                                SIRT1 impairs its ability ate cell senescence (By
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 391;
                                                                                                                                                                                             in a wide variety
ed or inactivated :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in a complex with
                                                                                                                                                                                                                                                                                                   for acetylation
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RESULT 9
P53_HUMAN

AC AC

P53 HUMAN STANDARD; PRT; 393 P04637; Q15086; Q15087; Q15088; Q16535; Q16810; Q16811; Q16848; Q86UG1; Q8J016;

393 AA

Q16807; Q99659;

Q16808; Q9BTM4;

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Pfam; PF07710; P53_Tetramer; 1
PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
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CHAIN 1 393 Cellular tumor antige
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30; GO:0005507; F:ATP binding; ISS.
30; GO:0005507; F:COPPET ion binding; ISS.
30; GO:0003677; F:DNA binding; ISS.
30; GO:0005515; F:DNA binding; ISS.
30; GO:0005515; F:protein binding; ISS.
30; GO:0005516; P:apoptosis; ISS.
30; GO:0006516; P:apoptosis; ISS.
30; GO:0006517; P:apoptosis; ISS.
30; GO:0006518; P:caspase-activation via
30; GO:0006518; P:caspase activation via
30; GO:0007050; P:cell aging; ISS.
30; GO:0007050; P:cell differentiation; ISS
30; GO:00030154; P:cell groliferation; ISS
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IPR011615; P53_DNA_bd.
IPR0112346; P53_RUNT_DNA_bd.
IPR010991; p53_tetramerīstn.
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P:cell cycle arrest; ISS.
P:cell differentiation; ISS.
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P:negative regulation of cell
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Interaction with HIPK2 (By simila
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N6-acetyllysine
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N6-acetyllysine
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"Isolation and of the human p5 EMBO J. 3:3257-
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                                                                                               Mirzayans R.;
"Isolation of two cell
                                                                                                                                              NUCLEOTIDE SEQUENCE, AND VARIANTS ARG-72 MEDLINE=93303270; PubMed=8316628;
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MEDLINE=85126934; PubMed=6396087;
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Rotter V.;
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Harris N., Brill E.,
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MEDLINE=85230577; PubMed=4006916;
Zakut-Houri R., Bienz-Tadmor B.,
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Name=TP53; Synonyms=P53;
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70:245-252(1988)
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3:3257-3262(1984).
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E., Shohat O., F
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PubMed-14660794; DOI=10.1073/pnas.29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted
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                                                                                             bombesin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Kidney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          restricted
                                                                                                                                                                                                                            NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Azuma K., Shichijo S., Itoh K.; "Identification of a tumor-rejection antigen recognized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Human TP53 from the malignant glioma-derived M059K has a cancer-associated mutation in exor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anderson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PubMed=11023613;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        J. Biol. Chem.
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                                                                                                           carcinoma
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muller E.H., Tilanus M.G.J.;
genomic sequence. Correctio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C.W.,
 Zhang A.H.;
the effect
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O.M., Almazov V.P.,

(JUN-1991) to the
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                                       SEQUENCE
                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                  of.
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\R-1997) to the EMBL/GenBank/DDBJ databases
                                                                                                           by anta
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necrosis factor cytotoxicity.";
276:3361-3370(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Allalunis-Turner M.J.;
                                                                                                       tant p53 expression and antagonists of growth l
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   of tumor
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                                                                          U.S.
                                       126-185
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EMBL/GenBank/DDBJ
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 suppressor
                                                                          100:15836-15841 (2003)
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                                                                                                             1 growth of DMS-15.
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extension

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RESULT 10
P53_MACFA
     SCORPTOR
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Best Local S
Matches 18
                                                                                                                                             P53_MACFA
P56423;
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Lipsky P.E.;
"Bcl6 and P5
                                                    15-JUL-1998, integrated into Unipr
10-MAY-2004, sequence version 2.
07-FBB-2006, entry version 50.
Cellular tumor antigen p53 (Tumor
Name=TP53; Synonyms=P53;
 Macaca fascicularis
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Thompson-Hehir J., Davies M.P.A., Green J.A., Halliwell N., Joyce K.A., Salisbury J., Sibson D.R., Vergote I., Walker C., "Mutation detection utilizing a novel PCR approach for amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nimri L.F., Owais W., Momani E.; "Detection of P53 gene mutations with cigarette smoking."; Submitted (AUG-2003) to the EMBL/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "The human tumour suppressor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=91141509; PubMed=1705009; Samad A., Carroll R.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pinto E.M., "Allelic var
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Ovarian adenocarcinoma; Filippini G., Soldati G.;
                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEAR LOCALIZATION SIGNAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      normal cells.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ALTERNATIVE SPLICING.
MEDLINE=96197761; PubMed=8632903;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      laman J.-M., Waridel F., Estreicher A., lbert D., Iggo R., Frebourg T.;
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                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ., Mendonca B.B., Latronico A.C.; variant in intron 9 of TP53 gene."; (APR-2003) to the EMBL/GenBank/DDBJ
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al cell, and Glial tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biol. 11:1598-1606(1991)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene from microdissected tissue:
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                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                              integrated into UniProtKB/Swiss-Prot
                                                                                                                                                                 STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mutations
                  (Crab eating macaque)
Chordata; Craniata; V
                                                                                                                                                                                                                                                                                                                                           94.3%;
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                                                                                                                                                                                                                                                                                                                                            Score 99;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene p53
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                                                                                                                                                                 PRT;
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                                                                        suppressor p53).
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       Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                            DB 1; L
. 2.9e-06;
                    e) (Cynomolgus monkey).Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vannier A., Limacher J.-M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       alternatively spliced
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       application to archival
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       R RNR; PS423; 94-289.

R GO; GO:0005739; C:mitochondrion; ISS.
R GO; GO:0005739; C:mitochondrion; ISS.
R GO; GO:0005739; C:mitochondrion; ISS.
R GO; GO:0005507; F:ATP binding; ISS.
R GO; GO:0005507; F:DNA binding; ISS.
R GO; GO:0005739; F:DNA strand annealing activity;
R GO; GO:0000739; F:DNA strand annealing activity;
R GO; GO:000515; F:protein binding; ISS.
R GO; GO:000515; P:apoptosis; ISS.
R GO; GO:0006204; P:base-excision repair; ISS.
R GO; GO:0006205; P:cell aging; ISS.
R GO; GO:0007569; P:cell aging; ISS.
R GO; GO:0007569; P:cell differentiation; ISS.
R GO; GO:0007509; P:cell proliferation; ISS.
R GO; GO:00008209; P:negative regulation of cell groups; ISS.
R GO; GO:0001088; P:negative regulation repair; ISS.
R GO; GO:0001089; P:negative regulation repair; ISS.
R GO; GO:0001089; P:negative regulation repair; ISS.
R GO; GO:0001080; P:negative regulation repair; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   replicative life span of macaque cells.";

Biophys. Res. Commun. 295:644-650(2002).

Biochem. Biophys. Res. Commun. 295:644-650(2002).

-:- FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    - -
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Khan M.A., Hansen C., Welsh J.A., Bennett W.P.
Submitted (FEB-1996) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                 EMBL; AF456343;
HSSP; P04637; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                      Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Copyrighted by the UniProt Consortium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Somatic mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=22094611;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cercopithecidae; Cercopithecinae; Macaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -!- SIMILARITY: Belongs to the p53 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COPACTOR: Binds 1 zinc ion per subunit (By similarity).

COBACTOR: Binds DNA as a homotetramer. Found in a complex with CABLES1 and p53/TP73. Interacts with histone acetyltransferases EP300 and methyltransferases HRWILL2 and CARMI. and recruits the to promoters. C-terminus interacts with TAF1, when TAF1 is part the TFIID complex. Interacts with HIPK1, HIPK2, AXIN1, and P53DINP1. P33DINP1. Part of a complex consistiting of TP53, HIPK2 and AXIN1. Interacts with WWOX (By similarity).

SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity).

PTM: Acetylated. Its deacetylation by SIRT1 impairs its ability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      by CREBBP (By similarity).
DISEASE: p53 is found in increased amounts in a wide variety transformed cells. p53 is frequently mutated or inactivated:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            which may prevent the interaction with MDM2. Phosphorylated on Thr-55 by TAF1 which promotes MDM2-mediated p53 degradation. Phosphorylated by HIPK1. Phosphorylated on Ser-46 by HIPK2 upon irradiation. Phosphorylation on Ser-46 is required for acetylation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PTM: Phosphorylated. Phosphorylation on Ser retranscriptional activation. Phosphorylated on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          many types of cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        induce proapoptotic program
                                                                                                                                                                                                                                                                                                                                                                                                  U48957; AAB91535.1;
   IPR002117; 1
IPR011615; 1
IPR012346; 1
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                                                                                                                                                                                                                                                                                                                                                                                   AAN64027.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PubMed=12099687; DOI=10.1016/S0006-291X(02)00730-1;
; P53.
; P53_DNA_bd.
; P53_RUNT_DNA_bd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      see http://www.uniprot.org/terms
Attribution-NoDerivs License
                                                                                                                                                                                                                                          activity;
                                                                                                                                                                   cytochrome
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                                                             growth; ISS ISS.
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on Thr-18 by VRK1,
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3_MACFU
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P61360;
10-MAY-2004, integrated into U
10-MAY-2004, sequence version
07-FEB-2006, entry version 25.
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Mammalia; Eutheria; Euarchontoglires; Primates;
Cercopithecidae; Cercopithecinae; Macaca.
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"Somatic mutations in the p53 gene account for the replicative life span of macaque cells."; Biochem. Biophys. Res. Commun. 295:644-650(2002).
                                            MEDLINE=22094611; Parishida
                                                                                                                                                                                                     Cellular tumor antigen p53
Name=TP53; Synonyms=P53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   METAL
METAL
                                                                           TISSUE=Lung;
                                                                                           NUCLEOTIDE SEQUENCE [MRNA].
                                                                                                                                                                                       Macaca fuscata fuscata (Japanese macaque)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRINTS; PR00386; P53SUPPRESSR. ProDom; PD002681; P53; 1. PROSITE; PS00348; P53; 1.
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Pfam; PF00870; P53; 1.
Pfam; PF07710; P53_tetramer;
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N6-acetyllysine
N6-acetyllysine
N6-acetyllysine
N -> H (in Ref. 1
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Pred. No.
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Cellular tumor antigen
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By similarity.
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                                                                                                                                                                                                                   suppressor
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                                                              DOI=10.1016/S0006-291X(02)00730-1;
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(By similarity).
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localization signal
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(by VRK1) (By
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 MOTIF
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                                                                                                                                                                                                                                                                                                                                                                          GO: 0005739
GO: 0005524
GO: 0005524
GO: 0005507
GO: 00007377
GO: 00005515
GO: 0006515
GO: 0006635
GO: 0007569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PTM: Phosphorylated on Thr-55 by TAF1 which p53 degradation (By similarity).
DISEAS: p53 is found in increased amounts i transformed cells. p53 is frequently mutated many types of cancer.
SIMILARITY: Belongs to the p53 family.
                                                                                                                                                                                                                                                                                                               IPR010991;
                                                                                                                                                                                                                                                                                                                           IPR002117; P53.
IPR011615; P53_DNA_bd.
IPR012346; P53_RUNT_DN
                                             102
1
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325
368
                             292
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321
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FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bcl-2
```

expression.

COFACTOR: Binds 1 zinc ion per subunit (By similarity).

SUBUNIT: Binds DNA as a homotetramer (By similarity). Found in a complex with CABLES1 and p53/TP73 (By similarity). Interacts with histone acetyltransferases EP300 and methyltransferases HRWT1L2 and CARMI, and recruits them to promoters. Interacts with WWOX Found in a steracts with (By

PTM: Phosphorylated on Thr-18 by VRK1, which interaction with MDM2 (By similarity). PTM: Acetylated. Its deacetylation by SIRT1 similarity).
SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity). which may prevent

induce proapoptotic program and modulate cell impairs its ability senescence (By ő

.n increased amounts in a wide variety
is frequently mutated or inactivated ; promotes MDM2-mediated

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AF456344; AAN64028.1; -; C:mitochondrion; C:nucleolus; ISS. mRNA. ISS.

7; F:NAA binding; ISS.
5; P:protein binding; ISS.
5; P:apoptosis; ISS.
4; P:base-excision repair; ISS.
5; P:call aging; ISS.
6; P:cell dying; ISS.
70; P:cell differentiation; ISS.
10; P:cell differentiation; ISS.
10; P:cell differentiation; ISS.
13; P:cell proliferation; ISS.
13; P:negative regulation of cell grow
19; P:neclectide-excision repair; ISS. F:ATP binding; ISS. F:copper ion binding; ISS cytochrome ů SSI

growth;

SSI

Pfam; PF00870; P53; 1. — Pfam; PF07710; P53 tetramer; 1. PR10710; P53 tetramers. ProDom; PD002681; P53; 1. PROSITE; PS00348; P53; 1. Acetylation; Activator; Anti-oncogene; Apoptosis; Cell cycle; DNA-binding; Metal-binding; Nuclear protein; Phosphorylation; Transcription regulation, Zinc.
1 393 Cellular tumor antigen
/FTId=PRO 0000185705.
By similarity. p53_tetrameristn. Interaction with HRMTIL2 (By similarity). Transcription activation (acidic). Interaction with WWOX (By similarity). Interaction with CARMI (By similarity). Oligomerization. Bipartite nuclear similarity). DNA bd (repression of DNA-binding) localization si (By similarity)

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P53

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RESULT 12
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ID P53_N
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Matches 18
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NUCLEOTIDE SEQUENCE (GENOMIC DNA).

Khan M.A., Hansen C., Welsh J.A., Bennett W.P.;

Submitted (FEB-1996) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis Induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bc1-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=94171042; PubMed=8125305; DOI=10.1016/0378-1119(94)90812-5; Kay H.D., Mountjoy C.P., Wu G., Cornish K.G., Smith L.J.; "Sequence of a cDNA encoding the p53 protein in rhesus monkey (Macaca mulatta).";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JUL-1998, sequence version 07-FEB-2006, entry version 50 Cellular tumor antigen p53 (TV Name=TP53; Synonyms=P53;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P56424;
15-JUL-1998, integrated into UniProtKB/Swiss-Prot.
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NCBI_TaxID=9544;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE [MRNA].
                                                                       SUBUNIT: Binds DNA as a homotetramer. Found in a complex with CABLES1 and p53/TP73. Interacts with histone acetyltransferases EP300 and methyltransferases HRMIL2 and CARM1, and recruits them to promoters. C-terminus interacts with TAF1, when TAF1 is part of the TFIID complex. Interacts with HIPK1, HIPK2, AXIN1, and P53DINP1. Part of a complex consisiting of TP53, HIPK2 and AXIN1. Interacts with WWOX (By similarity).
                                               SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity) PTM: Acetylated. Its deacetylation by SIRT1 impairs its abili-
                                                                                                                                                                                      COFACTOR: Binds 1 zinc ion per subunit (By similarity)
                             induce proapoptotic
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                              program
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zinc (By similarity).
Zinc (By similarity).
Zinc (By similarity).
Zinc (By similarity).
5'-phospho-RNA (covalent) (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          similarity).
N6-acetyllysine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Phosphothreonine (by TAF1) (By
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           similarity)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Phosphothreonine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Phosphoserine (by PRPK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ۳.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (By similarity).
(By similarity).
(By similarity).
(By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         suppressor
                                modulate cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Le . 2.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         γď)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 393;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         y similarity)
y similarity)
y similarity)
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                                senescence
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                                                 its ability to
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                                                                                                                                 of.
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   EMBL; L20442;
EMBL; U48956;
HSSP; P04637;
                                                                                                                                                                                                                                                                                              DNA BIND
REGION
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InterPro; IPR011615; P53 NA bd.
InterPro; IPR012346; P53 NNT DNA bd.
InterPro; IPR010991; P53 tetrameristn.
Pfam; PF00870; P53; 1.
Pfam; PF07710; P53 tetramer; 1.
PRINTS; PR00386; P53SUDPRESSR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SMR; P56424; 94-289.

GO; GO:0005739; C:mitochondrion; ISS.
GO; GO:0005730; C:nucleolus; ISS.
GO; GO:0005730; C:nucleolus; ISS.
GO; GO:0005524; F:ATP binding; ISS.
GO; GO:0005527; F:DAP binding; ISS.
GO; GO:000557; F:DAP binding; ISS.
GO; GO:0000577; F:DAP binding; ISS.
GO; GO:00005739; F:DAP binding; ISS.
GO; GO:00005215; F:protein binding; ISS.
GO; GO:00005215; F:protein binding; ISS.
GO; GO:00005284; P:base-excision repair; ISS.
GO; GO:00005284; P:base-excision via cytochrome c; ISG; GO:0000529; P:cell aging; ISS.
GO; GO:0000750; P:cell aging; ISS.
GO; GO:000750; P:cell differentiation; ISS.
GO; GO:0000753; P:cell proliferation; ISS.
GO; GO:0000203; P:negative regulation of cell growth; ISS.
GO; GO:0000529; P:nucleotide-excision repair; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOD_RES
                                                           MOD_RES
                                                                                                                                                 METAL
METAL
                                                                                                                                                                                                MOTIF
                                                                                                                                                                                                                                 MOTIF
                                                                                                                                                                                                                                                  REGION
REGION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transformed cells. p53 is frequently mutated many types of cancer.
-i- SIMILARITY: Belongs to the p53 family.
                                                                                                                                                                                 METAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Acetylation; Activator; Anti-oncogene; Apoptosis; Cell cycle; DNA-binding; Metal-binding; Nuclear protein; Phosphorylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ProDom; PD002681; P53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium,
                                                                                                                BINDING
                                                                                                                                   METAL
                                                                                                                                                                                                                                                                                                                                                                                                                                              Transcription;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     irradiation. Phosphorylation on Ser-46 is required the delignment of the CREBBP (By similarity).

DISEASE: p53 is found in increased amounts in a wide variety of piscase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transcriptional activation. Phosphorylated on Thr-18 by VRK1, which may prevent the interaction with MDM2. Phosphorylated on Thr-55 by TAF1 which promotes MDM2-mediated p53 degradation. Phosphorylated by HIPK1. Phosphorylated on Ser-46 by HIPK2 upon irradiation. Phosphorylation on Ser-46 is required for acetylati
                                                                                                                                                                                                                                                                 102
1
66
100
116
300
319
325
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; AAB91534.1;
; 1OLG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Transcription
                                                                                                                350
176
179
238
242
392
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                                                                                                                                                                                                                                                                             Transcription activation (acidic).
Interaction with WWOX (By similarity).
Interaction with HIPK1 (By similarity).
Interaction with AXIM1 (By similarity).
Interaction with CARM1 (By similarity).
Interaction with HIPK2 (By similarity).
                                                                                                                                                                                                                                                                                                                                                         By similarity.
Interaction with HRMT1L2 (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                regulation; Zin
Cellular tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mRNA.
                                                                                                                Zinc (By similarity).
Zinc (By similarity).
Zinc (By similarity).
5'-phospho-RNA (covalent)
                                                                                                                                                                                 Zinc
                                                                                                                                                                                                                  Bipartite r
similarity)
                              Phosphoserine
                                                   similarity)
                                                                Phosphoserine (by Phosphothreonine
                                                                                                                                                                                                  Nuclear export signal
                                                                                                                                                                                                                                                    Basic
                                                                                                                                                                                                                                                                   Oligomerization.
                                                                                                                                                                                                                                                                                                                                                                                                               Cellular tumor antigen /FTId=PRO_0000185706.
                                                                                                   similarity)
                                                                                                                                                                                 (By similarity)
                                                                                                                                                                                                                                                  (repression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             see
                                                                                                                                                                                                                                   nuclear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           http://www.uniprot.org/terms
                                   γď
                    (by TAF1)
                                                                PRPK) (By similarity) (by VRK1) (By
                                                                                                                                                                                                                                                    of.
                                     HIPK2) (By similarity)
                                                                                                                                                                                                                                   localization signal (By
                                                                                                                                                                                                                                                    DNA-binding)
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밁 á

(By similarity).

(Ву

p53

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RESULT 13
P53_TUPGB
ID P53_T
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Wild-type p53 sequence of tree shrews.";
Submitted (AUG-1999) to the EMEL/GenBank/DDBJ databases.
-i- FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOD_RES
MOD_RES
MOD_RES
SEQUENCE
HSSP; P04637; 1GZH.
SMR; Q9TTA1; 94-289.
GO; GO:0005739; C:mitochondrion;
GO; GO:0005730; C:nucleolus; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-2000, integrated int
01-MAY-2000, sequence versi
07-FEB-2006, entry version
                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE [MRNA].
STRAIN=Chinensis; TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Scandentia; Tupaiidae; Tupaia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-FEB-2006, entry version 50.
Cellular tumor antigen p53 (Tumor suppressor p53)
                                                                         EMBL; AF175893; AAF22640.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tupaia glis belangeri (Common tree shrew)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Name≖TP53; Synonyms≂P53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q9TTA1;
                                                                                                                                                                                                                                                                                                                                                                                                 COPACTOR: Binds 1 zinc ion per subunit (By similarity).

SUBUNIT: Binds DNA as a homotetramer. Interacts with histone acetyltransferases EP300 and methyltransferases HRWT12 and CARM1, and recruits them to promoters. C-terminus interacts with TAF1, when TAF1 is part of the TFIID complex. Interacts with HIPK1, when TAF1 is part of the TFIID complex consisting of TP53, HIPK2, AXIN1, and P53DINP1. Part of a complex consisting of TP53, HIPK2, HIPK2, CONSTRUCT WITH WMOX (By similarity).

SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity).

SUBCELLULAR LOCATION: Cytoplasmic and nuclear (By similarity).

PTM: Acetylated. Its deacetylation by SIRT1 impairs its ability to
                                                                                                                                                               transformed cells. p53 is frequently mutated or inactivated many types of cancer.
SIMILARITY: Belongs to the p53 family.
                                                                                                                                                                                                                  transcriptional activation. Phosphorylated on Thr-18 by VRKI, which may prevent the interaction with MDM2. Phosphorylated on Thr-55 by TAR1 which promotes MDM3-mediated p53 degradation. Phosphorylated by HIPK1 (By similarity). Phosphorylated on Se by HIPK2 upon UV irradiation. Phosphorylation on Ser-46 is required for acetylation by CREBBP (By similarity). DISEASE: p53 is found in increased amounts in a wide variety.
                                                                                                                                                                                                                                                                                                                                                 PTM: Phosphorylated. Phosphorylation on Ser residues mediates
                                                                                                                                                                                                                                                                                                                                                                      similarity)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression
                                                                                                                                                                                                                                                                                                                                                                                   induce proapoptotic program
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence version 1.
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373
382
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ¥
                                                                         -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N6-acetyllysine (By
N6-acetyllysine (By
N6-acetyllysine (By
N6-acetyllysine (By
E212E5E4FE650103 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                       modulate cell
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(By similarity).
(By similarity).
(By similarity).
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                                                                                                                                                                                                                                                                                                           hr-18 by VRK1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                     senescence
                                                                                                                                                                                                                                                          on Ser-46
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Best Local S
Matches 18
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SEQUENCE
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                                                                                                                                                                                             METAL
BINDING
                                                                                                                                                                                                                                     MOTIF
METAL
                                                                                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF00870; P53; 1.
Pfam; PF07710; P53 tetramer; 1.
PRINTS; PR00186; P53sUPPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro;
InterPro;
                                                                                                                                                           MOD_RES
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                                                                                                                                                                                                                 METAL
                                                                                                                                                                                                                            METAL
                                                                                                                                                                                                                                                                                                                                                                                                      Transcription;
                                                                                                                                                                                                                                                                                                                                                                                                              Acetylation; Activator; Anti-oncogene; Apoptosis; DNA-binding; Metal-binding; Nuclear protein; Phosp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GO: 0005524
GO: 0003577
GO: 0003577
GO: 0000739
GO: 0006515
GO: 0006284
GO: 0007569
GO: 0007050
12 PPLSQETFSDLWKLLPEN
                  1 PPLSQETFSDLWKLLPEN 18
                                       18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               :0006289;
                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IPR002117; P53_DNA_bd.
IPR011615; P53_DNA_bd.
IPR012346; P53_RUNT_DNA_bd.
IPR010991; p53_tetramerIstn.
                                        Conservative
                                                                                                                                 55
                                                                               AA;
                                                                                                                                                                                                                                                                                                                                                                      Transcription regulation; Zinc.

Cellular tumor antigen p53

/FTId=PRO 0000185715.

2 292 By similarity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P:cell proliferation; ISS.
P:negative regulation of cell
P:nucleotide-excision repair;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F:ATP binding; ISS.
F:copper ion binding; ISS
F:DNA binding; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P:cell aging; ISS.
P:cell cycle arrest; ISS.
P:cell differentiation; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P:base-excision repair;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             F:DNA strand annealing activity; F:protein binding; ISS.
P:apoptosis; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P:caspase activation via cytochrome
                                                                                                                                 55
                                                                               43552
                                                                                                                                                              18
                                                94.3%;
                                                 .0%;
                                                                               ¥.
29
                                      0
                                      Score 99; DB; Pred. No. 2.9
                                                                                                                           similarity).

Phosphoserine (by HIPK2) (By similarity)

Phosphorhreonine (by TAF1) (By
                                                                                        N6-acetyllysine
N6-acetyllysine
                                                                                                                                                                                                                            Zinc
                                                                                                                                                                                                                                                                                             Transcription activation (acidic). Interaction with WWOX (By similarity). Interaction with HIPK1 (By similarity). Interaction with AXINI (By similarity). Interaction with CARMI (By similarity). Interaction with HIPK2 (By similarity).
                                                                                                                                                                                           Zinc (By similarity).
5'-phospho-RNA (covalent)
                                                                                                                                                                                                                                     Nuclear export signal Zinc (By similarity).
                                                                                                                                                                                                                                                                   Basic (repression 
Bipartite nuclear
                                                                                                           N6-acetyllysine
                                                                                                                      similarity).
                                                                                                                                                              Phosphothreonine
                                                                                                                                                                       Phosphoserine (by
                                                                                                                                                                                   similarity)
                                                                                                                                                                                                                                                                                        Oligomerization.
                                                                                                                                                                                                                                                                                                                                                             Interaction with HRMT1L2
                                                                               FD9F36603945A1FA
                                                                                                                                                                                                                            (By similarity)
                                                                                                                                                                                                                 (By similarity)
                                   DB 1; L. 2.9e-06; 0;
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                                                                                       (By similarity).
(By similarity).
(By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                              osis; Cell cycle;
Phosphorylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              growth; ISS ISS.
                                                                                                                                                            (by VRK1)
                                                        Length 393
                                                                                                                                                                                                                                                                  of DNA-binding).
localization signal
                                                                                                                                                                        PRPK)
                                                                               CRC64;
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                                                                                                                                                                                                                                               (By similarity).
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                                                                                                                                                                        similarity).
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                                       Gaps
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RESULT 14
Q5U0E4 HUMAN
D Q5U0E4;

REC Q5U0E4;

DT 07-DEC-2004, integrated into UniProtKB/TrEMBL
DT 07-DEC-2004, sequence version 1.

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RESULT 15
QZXN98 HUMAN
ID QZXN98 H
AC QZXN98;
DT 20-DEC-2
DT 20-DEC-2
DT 20-MAR-2
DT 97-MAR-2
DE P53 prot
GN Name=TP5
QN Homo sap
QC Eukaryot
QC Mammalia
QC Homo.
QX NCBI_Tay
RN [1]
RX PUBMed=1]
RX PUBMed=1
RA Bourdon
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR002117; P
InterPro; IPR011615; P
InterPro; IPR012346; P
InterPro; IPR010991; p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SMR; Q500
Ensembl;
G0; G0:00
                                                                                                                                                             Q2XN98;
20-DEC-2005, integrated into UniProtKB/TrEMBL.
20-DEC-2005, sequence version 1.
07-MAR-2006, entry version 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pfam; PF00870; P53; 1.
Pfam; PF07710; P53 tetramer; 1.
PRINTS; PR00386; P53sUPPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-FEB-2006, entry version 10.
Tumor protein p53 (Li-Fraumeni syndrome).
Homo sapiens (Human).
TISSUE=Normal colon;
PubMed=16131611; DOI=10.1101/gad.1339905;
Bourdon J.C., Fernandes K., Murray-Zmijewski F.,
                                                                                                  Bukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                              Activator, Anti-oncogene, Apoptosis, Cell cycle, Metal-binding, Nuclear protein, Phosphorylation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Koundinya M., Raphael J., Moreira D., Kelley T., LaB Phelan M., Farmer A.; "Cloning of human full-length CDSs in BD Creator(TM)
                                                                                                                                                                                                                         QZXN98_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                    Transcription regulation; Zinc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (OCT-2004) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
Kalnine N., Chen X.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota;
                                               NUCLEOTIDE SEQUENCE.
                                                                    NCBI_TaxID=9606;
                                                                                                                              Homo sapiens (Human)
                                                                                                                                             Name=TP53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GO:0005634; C:nucleus; IEA.
GO:00046872; F:metal ion binding; IEA.
GO:0003700; F:transcription factor activity; IEA.
GO:0008270; F:zinc ion binding; IEA.
GO:0008270; F:zinc ion binding; IEA.
GO:0006915; P:apoptosis; IEA.
GO:0007049; P:cell cycle; IEA.
GO:0007049; P:cell cycle; IEA.
GO:0006356; P:regulation of progression through . . . .
GO:0006350; P:transcription; IEA.
                                                                                                                                                         protein.
                                                                                                                                                                                                                                                                                              12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BT019622; AAV38428.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                        μ
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                                                                                                                                                                                                                                                                                             PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENSG00000141510; Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                          43730 MW;
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                                                                                              Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P53. DNA_bd.
P53_RUNT_DNA_bd.
P53_RUNT_DNA_bd.
P53_tetrameristn.
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                                                                                                                                                                                                                                                                                                                                                                             94.3%;
                                                                                                                                                                                                                                                                                                                                                               100.0%;
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                                                                                                                                                                                                                            393 AA.
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       Liu
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Transcription;
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kaer J., Lin
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       Diot
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       Α.
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SKWARRCCCCRFA
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Best Local S
Matches 18
                                                                                                            EMBL; DQ286964; ABB80266.1; -; mRNA.
EMBL; DQ191317; ABB80262.1; -; mRNA.
Activator; Anti-oncogene; Apoteosis; Cell cycle; DNA-binding; Metal-binding; Notein; Phosphorylation; Transcription; Transcription; Zinc.
SEQUENCE 393 AA; 43712 MW; AC611E4938C7EC3B CRC64;
                                                                                                                                                                                                             Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                   "p53 isoforms can regulate p53
Genes Dev. 19:2122-2137(2005).
                                                                                                                                                                                                                                                                                Xirodimas D.P.,
12
              1 PPLSQETFSDLWKLLPEN
                                                      l Similarity
18; Conser
  PPLSQETFSDLWKLLPEN
                                                    94.3%;
ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                  Saville M.K.,
                           18
                                                       ,
  29
                                                                    Score 99; pred. No.
                                                                                                                                                                                                                                                                                  Lane D.P.;
                                                                                                                                                                                                                                                                   transcriptional activity.";
                                                         Mismatches
                                                                    DB 2; Lo
                                                                                   Length 393;
                                                         Indels
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Search completed: July 5, 2006, 22:45:46
Job time : 296 secs

Copyright

GenCore version (c) 1993 - 2006

5.1.9 Biocceleration Ltd

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Minimum
Maximum
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Maximum Match 100%
Listing first 45 s
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Perfect score:
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1: geneseqp198
2: geneseqp199
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105
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            geneseqp2003bs:*
geneseqp2004s:*
geneseqp2005s:*
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geneseqp2002s:*
geneseqp2003as:*
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44.322 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	8	7	თ	σ	4.	ω	N	_	Result No.
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71	68	64	64	64	64	64	64	64	52	50	41	36	32	32	32	26	25	25	20	18	19	19	Length D
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AAW47079	ADB71206	AAW94303	AAW42970	AAW42878	AAW57240	AAW48243	AAW07886	AAR42174	AAW13602	ADT02859	AAW13603	AAW13604	ADZ48535	ADJ25788	AAB30523	AAW60202	AAR51879	AAR54907	AAE30863	AAW37228	AAW82319	AAW82321	ID
Aaw47079 Human p53	Adb71206 Human p53	Aaw94303 Human MDM	Aaw42970 N-termina	Aaw42878 N-termina	Aaw57240 Human p53	Aaw48243 Human p53	Aaw07886 Human p53		Aaw13602 p53 prote		Aaw13603 p53 prote	Aaw13604 p53 prote	Adz48535 Human p53	Adj25788 MDM2 bind	Pep	Aaw60202 p53 pepti	Aar51879 Human p53	Aar54907 Immunodom	Aae30863 EGFP-BOX-	Aaw37228 p53 N-ter	Aaw82319 p53 homol	Aaw82321 p53 homol	

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24
99	99	99	99	99	99	99	99	99	99	99	99	99	99	99	99	99	99	99	99	99	99
94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3
353	343	341	337	293	283	261	260	241	225	211	164	161	73	73	73	73	73	73	73	73	73
6	7	œ	N	7	4.	w	4	N	4.	4	.Δ	4.	10	10	10	10	10	10	10	10	10
AAE35350	AAE39685	ADN34522	AAW13962	AAE39686	ABG01512	AAY70714	ABG03222	AAR51872	AAB36690	AAB36685	AAB36689	AAB36683	AEE18140	AEE18139	AEE18144	AEE18137	AEE18125	AEE18138	AEE18143	AEE18142	AEE18141
Human	Aae39685 Human p53	Adn34522 Human p53	Aaw13962 Chimeric	Aae39686 Human p53	Abg01512 Novel hum	Aay70714 Human p35	Novel	Aar51872 Human p53	Aab36690 Mammalian	Aab36685 Mammalian	Aab36689 Mammalian	Aab36683 Mammalian	Aee18140 Human p53	Aee18139 Human p53	Aee18144 Human p53	Aee18137 Human p53	Aee18125 Human p53	Aee18138 Human p53	Aee18143 Human p53	Aee18142 Human p53	Aee18141 Human p53

ALIGNMENTS

RESULT 1 AAW82321 ID AAW8

AAW82321 standard; peptide; 19

⋛

29-OCT-1998. Synthetic. p53 homologue TIP peptide. (UYDU-) UNIV DUNDEE 22-APR-1997; 20-APR-1998; WO9847919-A1 p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament. 22-FEB-1999 AAW82321; (first entry) 97GB-00008089 98WO-GB001140.

New substance with a mdm2 binding domain and coupling partner - useful for stabilising in cells without an efficient mdm2-mediated degradation

WPI; 1998-609975/51.

Lane DP;

Disclosure; Fig 1; 52pp; English.

This sequence is a peptide homologue of a region of p53 which binds to mdm2. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows level p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 function. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53

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RESULT 2
AAW823119
ID AAW8
AX AAW8
AX AAW8
AX AAW8
AX P53;
XX Synt
XX Synt
XX WO9E
XX WO9E
XX OS Synt
XX WO9E
XX U91
XX WO9E
XX WO9E
XX U91
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                                                     AAW37228
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Matches 19
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Best Local
                                                                                                                                                                                                                                                                                                                          This sequence is a peptide homologue of a region of p53 which binds to mdm2. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 function. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New agents which inhibit interaction of p53 and mdm2 - useful activating p53, e.g. for treating cancers, viral conditions of conditions associated with non functional p53 or mdm2.
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 AAW37228;
                                  AAW37228 standard; peptide; 18 AA
                                                                                                                                                                                                                                                                          Sequence 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW82319 standard;
                                                                                                                                                                                                                                                                                                               non-functional p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 1; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-609932/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lane DP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19;
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                                                                                                                                                                                                                   100.0%; Score 105; DB 2; 100.0%; Pred. No. 1.7e-09;
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Pred. No. 1.7e-09;
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 RESULT 4
AAE30863
ID AAE3
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AC AAE3
AC AAE3
DT 24-E
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DE EGFF
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Garcia-Echeverria C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-1996;
07-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumour;
                   EGFP-BOX-1 domain phospho-peptide used
                                                       24-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9801467-A2
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                                                                                                                                                                                                                                                                                                                  Local
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binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2/p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia patients and for treatment or prevention of disease involving p53/MDM2 interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and accumulation in normal cells. The peptides which mimic the MDM2 binding site in p53, have a significantly greater blocking activity compared with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            assay for analysing the interaction between human oncogenic protein MDI and p53. The invention provides peptide derivatives capable of binding the human MDM2. These peptides can specifically inhibit or block the binding the human MDM2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compounds binding to MDM2 protein and inhibit its useful in, e.g. diagnosis and treatment of cancer and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This represents a p53 N-terminal peptide fragment used in an Elisa TIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 34; 45pp; English.
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                                                                                                                                                                 AAE30863 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oncogenic protein; p53; r; diagnosis; binding;
                                                                                                                                                                                                                                                                                                                                     1 PPLSQETFSDLWKLLPEN 18
                                                  (first entry)
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97GB-00007041.
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                                                                                                                                                                                                                                                                                                                                                                                                                94.3%; but
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Boettger
Chene P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         p53; human; inhibition; interaction; cancer;
ng; viral infection; Elisa TIP assay.
                                                                                                                                                                                                                                                                                                                18
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Pred. No.
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Furet P;
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in the invention
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RESULT 5
AAR54907
ID AAR5
XX AAR5
AC AAR5
XC AAR5
XC Immu
XX Immu
XX Canc
KW immu
KW trar
XX Homc
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                 Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a peptide for use in modulating the binding of p53 polypeptide to a p300 polypeptide. The new peptide is useful in modulating the binding of a p53 polypeptide to a p300 polypeptide. The peptide may be used to regulate the mammalian cell cycle or to induce or prevent cell death, for the treatment of cancer or ischaemia. The invention is useful in gene therapy. The present sequence is EGFP-BOX-1 domain phospho-peptide used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                               transformation-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                   cancer: pre-cancerous state; detection; diagnosis; human p53
immunodominant epitope; human cellular tumour antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
29-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New peptide for modulating the binding polypeptide, useful for regulating the treatment of cancer or ischemia.
                         Soussi T,
                                                                                                                                                                  02-NOV-1993;
                                                                                                                                                                                                                11-MAY-1994.
                                                                                                                                                                                                                                                                 WO9410306-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunodominant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR54907;
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                                                                       (EURO-) LAB EUROBIO
                                                                                                                     02-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 22; 87pp; English.
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                      Lubin R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                          epitope from p53 N-terminal
                                                                                                                     92FR-00013110
                                                                                                                                                                     93WO-FR001082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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Pred. No.
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J. 1.7e-08;
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RESULT 6
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Best Local S
Matches 18
                                Antibodies specific for p53 are detected by binding to immobilised fragments of the p53 gene product containing the antibody-binding region. Preferred fragments contain amino acids 1-241, 40-349, 40-393, 66-241, 66-393, 237-349, 237-393 and esp. 9-33, 37-52 or 368-386. See AARS1872-R51881 for sequences of these fragments. (Updated on 25-MAR-2003 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically react with anti-p53 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (AAR54907-R54921) are useful for
                                                                                                                                  Non-radioactive detection of p53 specific antibodies - by capture immobilised p53 or its fragments, then reaction with labelled seccantibody, for diagnosis of tumours and suitable for screening.
                                                                                                                                                                                                                                                                                                                                                                                                   Human nuclear phosphoprotein p53; tumour suppressor gene product; anti-oncogene; cancer; tumour; antibody binding region; epitope.
Sequence
                                                                                                           Claim 11; Page 19; 35pp; German
                                                                                                                                                                                                  WPI; 1994-135732/16.
                                                                                                                                                                                                                          Zentgraf H,
                                                                                                                                                                                                                                                                          30-SEP-1992;
                                                                                                                                                                                                                                                                                                   30-SEP-1993;
                                                                                                                                                                                                                                                                                                                            14-APR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
18-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR51879 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detecting and monitoring cancerous and precancerous conditions. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                     WO9408241-A1
                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR51879;
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                                                                                                                                                                                                                                                   (DEKR-)
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(first ent
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100.0%; Pr
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2.1e-08;
hes 0;
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Query Match Best Local Similarity

94.3%;

Score Pred.

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DB 2; L 2.1e-08;

Length

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RNA processing;

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RESULT 7
AAW60202
ID AAW6
XX AAW6
AC AAW6
XX AAW6
XX AAW6
XX AAW6
XX P53
XX Huma
XX P53
XX P03
XX P07-(
XX O1-(
XX O1-
RESULT 8
AAB30523
ID AAB3
XX
AC AAB3
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DT 06-M
XX
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptides AAM60202-05 are derived from human p53 protein. The present peptide corresponds to residues 11-35. The pptides are used in the method of the invention. Antibodies that bind to p53 protein are detected by immobilising a p53 peptide directly on to a flow cell of a sensor chip in a biosensor, treating the peptide with a sample of patient serum, diluted in buffer, and measuring binding of antibody to the peptide using the biosensor. The method is used to monitor cancer patients undergoing p53 gene therapy (to determine if an immune response has developed), and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting antibodies that bind p53 by reaction with immobilised p53 peptide(s) - attached directly to flow cells in the sensor chip of biosensor, used to analyse serum from cancer patients, e.g. those being given p53 gene therapy.
                          06-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 26 AA;
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                                                                        AAB30523;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                p53 peptide used to detect antibodies against p53
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                                                                                                                 AAB30523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 also to detect antibodies against mutant forms of p53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; p53; antibody; detection; biosensor; cancer patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DT,
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                                                                                                                   standard;
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                                                                                                                 peptide;
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2.2e-08;
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                                                                                                                 RESULT 9
ADJ25788
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The specification describes a method of synthesis of biologically active CC substances of determined structure directly in the cells of living CC organisms containing specific RNA or DNA sequence. The method is based on CC the hybridisation of two or more oligomers bound with biologically CC inactive substances to specific RNA or DNA in vivo in the cells of living CC organisms. After hybridisation of the oligomers, the biologically CC inactive precursors bound to the oligomers can interact with each other CC to make the active form of the substances. This changing of properties is CC due to chemical reactions which bind the biologically inactive precursors CC through a chemical bond into a biologically active form of the whole compound. The methods are useful for producing biologically active compounds from inactive precursors. These compounds may be inhibitors or Stimulators of cellular metabolism, DNA replication, RNA transcription, CC RNA translation, RNA elongation RNA processing, protein synthesis, CC protein processing, cellular fifterentiation, cell division, ion channel CC transmission, cellular protein and RNA transportation, processes of CC cellular oxidation, toxins, proteins or RNAs. AAB30523-36 represent CC peptides are fragments of the two compounds activator mRNA (AAC62167-60) or human NbHOT (AAC62181-94). The method of the invention is used to cardiot the two componers and
                                                                                                                                                                                                                                                                                                    Matches
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    20-MAY-2004
                                                                                           ADJ25788 standard; peptide; 32
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    producing proteins or polynucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel methods for the synthesis of biologically active compounds inactive precursors in the cells of living organisms, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-006911/01.
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                                                ADJ25788;
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                                                                                                                                                                                                                                                                                               94.3%; bu
100.0%; Pr
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                                                                                                                                                                                                                                                                                                                              Score 99;
Pred. No.
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Matches 18
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31-OCT-1997;
31-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of identifying a ligand that can mediate the biological activity of target protein via inhibition of the binding of target protein to a binding partner ligand comprising screening first combinatorial library having first member ligands for binding to target protein to identify target-binding ligand(s). The method is useful for identifying ligands that can mediate the biological activity of target proteins via inhibition of the binding of target proteins to a binding partner ligand. The invention does not require that the natural binding partner be used as reagent. The need for the natural binding partner be used as reagent. The need for the natural binding partner be used as reagent.
                                                                                                                    Human p53
                                                                                                                                                                                                                                                                                                                                                                               Sequence 32
              31-MAR-2005
                                        US2005069951-A1
                                                                                                                                               16-JUN-2005
                                                                                                                                                                                                 ADZ48535 standard; peptide; 32 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identification of ligand that can mediate biological activity of target protein, comprises screening first combinatorial library having first member ligands for binding to target protein to identify target-binding
                                                                                        combinatorial library; peptide library; p53
                                                                                                                                                                                                                                                                                                                                                                                                       present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; SEQ ID NO 48; 98pp; English.
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                                                                                                                    peptide SEQ ID NO:48
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97WO-US019638.
98US-00050359.
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Pred. No.
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                                                                                                                           CC first library, obtaining ligands which inhibit the binding of target color of a second library, and determining inhibitory ligands mediating activity cof the target protein. Also described is a method for identifying a pair cC of higands, the first ligand of the pair binding to a target protein, and cC the second library comprising several first member ligand cC the target protein, which involves: (a) screening the first ligand cC to the target protein, which involves: (a) screening the first ligand cC combinatorial library comprising several first member ligands for binding CC combinatorial gands; the pair inhibiting the binding of the first ligand cC member ligands for the ability to inhibit the binding several second cC member ligands for the ability to inhibit the binding of one or more carget binding gigands; where the second library is not identical to the first library, the first ligands of the first library is not identical to the first library, the first ligands of the second library and when the cfirst library, the second library, and when the gigand of the pair cc ligands are peptides, they are not more than 41 amino acids long, or they cc ligand which can mediate the binding of a target protein can be one associated with human cytomesus protein hibbition of the binding of a target protein kinase (preferably human protein kinase (preferably human protein kinase ce preferably human protein kinase ce peticals, where the ligand identified by (M1), is useful for crompunds which mediate the ligand identified by (M1), is useful for compunesses related to aberrant expression of the above target protein.

CC ligands which mediate the activity through binding to the receptor are contained by man protein and contained the binding of a target protein, even when the contained contained the activity through binding to the receptor are contained by man protein and contained the activity through binding to the receptor are contained by man protein are contained to the protein are contained to the protein are con
                                        Query Match
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Matches 18
                                                                                                                      Sequence 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method (M1) for identifying a ligand which can mediate the biological activity of a target protein via inhibition of the binding of a target protein to a binding partner. The method comprises identifying target-binding ligands by screening first member ligands of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying ligand mediating activity of target protein, by screening ligand libraries to obtain target-binding ligands and ligands inhibiting binding of target-binding ligand to target, determining ligands inhibiting activity of target.
                                                                                                                                                               not already known. which is used in a
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30-APR-1998;
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31-OCT-1997;
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1 PPLSQETFSDLWKLLPEN 18
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HYDE-DERUYSCHER
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KAY B K.
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                                          Conservative
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97WO-US019638.
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                                          Mismatches
                                                                                                                                                               present
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                                                             DB 9; L
                                                                                Length 32;
                                          Indels
                                          <u>,,</u>
                                          Gaps
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AAM13604

AAM136
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AAW13603
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The peptides AAW13602-6 represent peptide fragments derived from the wild type human p53 protein. This peptide corresponds to amino acids 6-41 of the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the murine double minute-2 (mdm2) protein (AAW13600). Mdm2 protein is a 90 kD phosphoprotein which has now been shown that the mdm2 protein tesulf has oncognic properties, especially in a p53-null background. Mdm2 is observed to unblock cell cycle arrest in G1 caused by over-expression of the p107 protein. The p53 CC peptides are examples of antagonists of the invention which are able to inhibit the oncogenic activity of mdm2. The antagonists are used to treat e.g. adenocarcinoma of the colon; cancer of the breast, lung or stomach; myeloid leukaemia; B cell lymphoma, or other hyperproliferative conditions such as restenosis. Note: this sequence is not given in the colon; cancer of the breast of the man p53 sequence
                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JAN-1998
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                          16-JAN-1998
                                                                                   AAW13603;
                                                                                                                                     AAW13603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page; 43pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treating cancer with antagonist of oncogenic activity of protein Mdm2 or nucleic acid encoding an antagonist, also viral vectors contg. this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-192837/17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9709343-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-MAR-1997
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                                                                                                                                                                                                                                                                                                                                                                                               18;
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INST NAT SANTE & RECH M
                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                        standard;
                                                                                                                                                                                                                                                                              PPLSQETFSDLWKLLPEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dubs-Poterszman M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A.
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                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                          (first entry)
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                                                                                                                                  peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%;
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                                                                                                                                        A
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3.2e-08;

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JP2004231535-A

Homo sapiens

Human p53 protein

fragment,

target of

an

immunoassay SeqID

0;

immunoassay; autoantibody; p53; human.

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RESULT 13
ADT02859
ID ADT02
XX
AC ADT02
XX
AC ADT02
XX
DT 02-DE
XX
Human
XX
KW immur
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OS Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                         The peptides AAW13602-6 represent peptide fragments derived from the wild C type human p53 protein. This peptide corresponds to amino acids 1-41 of CC the p53 sequence. The peptides are claimed peptides which are able to CC bind the N-terminal amino acids (1-134) of the murine double minute-2 (mdm2) protein (AAW1360). Mdm2 protein is a 90 kD phosphoprotein which CC binds and modulates the activity of the tumour suppressor protein p53. It CC has now been shown that the mdm2 protein itself has oncogenic properties, CC especially in a p53-null background. Mdm2 is observed to unblock cell CC cycle arrest in G1 caused by over-expression of the p107 protein. The p53 CC peptides are examples of antagonists of the invention which are able to CC inhibit the oncogenic activity of mdm2. The antagonists are used to treat CC e.g. adenocarcinoma of the colon, cancer of the breast, lung or stomach; CC myeloid leukaemia; B cell lymphoma, or other hyperproliferative conditions such as restenosis. Note: this sequence is not given in the CC specification but is constructed from the wild type human p53 sequence
                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse; Mdm2; murine double minute; phosphoprotein; binding; modulation; tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breast lung; stomach; myeloid leukaemia; lymphoma; hyperproliferative;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Treating cancer with antagonist of oncogenic activity of protein \mathsf{Mdm2} or nucleic acid encoding an antagonist, also viral vectors contg. this
                                                                                                                                02-DEC-2004
                                                                                                                                                                 ADT02859;
                                                                                                                                                                                                 ADT02859 standard;
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 41 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               restenosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RHON ) RHONE POULENC RORER SA.
                                                                                                                                                                                                                                                                                  12
                                                                                                                                                                                                                                                                                                                 1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                  PPLSQETFSDLWKLLPEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dubs-Poterszman M,
                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                (first entry)
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                                                                                                                                                                                                protein;
                                                                                                                                                                                                                                                                                                                                                                 94.3%;
                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                 Score 99;
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                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                   Indels
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RESULT 14
AAW13602
ID AAW13
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a novel reagent used for performing an immunoassay for detecting autoantibodies with respect to the p53 gene product. Specifically, it refers to a sandwich method that involves assaying an anti-human immunoglobulin antibody to determine the presence or absence of an autoantibody with respect to the p53 gene product in a test substance (such as blood) and comparing these values with a standard control from a healthy subject. The present invention describes a method based on a chemiluminescent enzyme immunoassay. As such, it provides an accurate and sensitive assay that can be used to detect and diagnose a second control from the present invention describes and accurate and sensitive assay that can be used to detect and diagnose a
                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                 Mouse; Mdm2; murine double minute; phosphoprotein; binding; modulation; tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breastlung; stomach; myeloid leukaemia; lymphoma; hyperproliferative;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reagent for performing immunoassay for detecting autoantibody against p53 gene product, comprises amino acid sequence, in which 8-50 amino acids are essential structural unit.
                                                                                                                                        02-SEP-1996;
                                                                                                                                                                                                                                                                                                                                        p53 protein
                                                                                                                                                                                                                                                                                                                                                                      16-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                    AAW13602
                                                                                                                                                                                                                                                                                                                                                                                                                              AAW13602 standard; peptide; 52 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                      Tocque B,
                                                                                                             04-SEP-1995;
                                                                                                                                                                    13-MAR-1997
                                                                                                                                                                                                WO9709343-A2
                                                                                                                                                                                                                                                        restenosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention.
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                                                                   RHONE POULENC RORER SA.
INST NAT SANTE & RECH N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 1; 91pp; Japanese.
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                                      Dubs-Poterszman
                                                                                                                                                                                                                                                                                                                                           amino acids 1-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA;
                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This
                                                                                                             95FR-00010331.
                                                                                                                                        96WO-FR001340
                                                                 SANTE & RECH MEDICALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polypeptide sequence is a human p53 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94.3%;
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Pred. No. 4.5e-08;
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                                                                                                                                                                                                                                                                                 antagonist;
ancer; breast;
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RESULT 15
AAR42174
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the murine double minute-2 (mdm2) protein (AAW13600). Mdm2 protein is a 90 kD phosphoprotein which binds and modulates the activity of the tumour suppressor protein p53. It has now been shown that the mdm2 protein itself has oncogenic properties, especially in a p53-null background. Mdm2 is observed to unblock cell cycle arrest in G1 caused by over-expression of the p107 protein. The p53 peptides are examples of antagonists of the invention which are able to inhibit the oncogenic activity of mdm2. The antagonists are used to treat e.g. adenocarcinoma of the colon; cancer of the breast, lung or stomach; myeloid leukhaemia; B cell lymphoma, or other hyperproliferative conditions such as restenosis. Note: this sequence is not given in the specification but is constructed from the wild type human p53 sequence
         Diagnosing neoplasia from amplification expression, also new DNA, MDM2 protein, sarcoma by inhibiting MDM2 expression.
                                                                                                                                                           07-APR-1992;
23-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                   p53 gene; tumour suppressor gene; regulation; cellular proliferation; cellular transformation; carcinoma; human; tumour; MDM2; inhibition;
                                                                                                                                                                                                       07-APR-1993;
                                                                                                                                                                                                                                    14-OCT-1993.
                                                                                                                                                                                                                                                                                                            Binding-site
                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                p53 N-terminal fragment
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05-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR42174 standard; peptide; 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 52 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page; 43pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating cancer with antagonist of oncogenic activity of protein Mdm2 or nucleic acid encoding an antagonist, also viral vectors contg. this
                                                                     WPI; 1993-336944/42.
                                                                                                   Burrell
                                                                                                                                                                                                                                                                  WO9320238-A2
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                                                                                                                               (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                     amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
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                                                                                                 Hill DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                           92US-00867840
92US-00903103
                                                                                                                                                                                                       93WO-US003199
                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                              /note=
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100.0%; Pred. No.
                                                                                                 Kinzler
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                                                                                                 Vogelstein
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                           of MDM2 gene - or elevate
antibodies and treatment
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                                        or elevated
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Search completed: July 5, 2006, 22:40:46 Job time: 198 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This sequence represents the product of the p53 gene. p53 is a tumour suppressor gene and this protein appears to be a member of a group of proteins which regulate normal cellular proliferation and suppression of cellular transformation. Inactivation of the p53 gene has been implicated in the formation, or progression of a wide variety of carcinoma. p53 is thought to interact with the protein product of a gene which is amplified in a number of human tumours, designated MDM2. Polypeptides containing at least amino acids 13-41 of p53, or the DNA encoding these, may be used to inhibit the growth of tumour cells containing MDM2 gene amplification. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 35; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 64 AA;
                                                                                                                                                     12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                1 PPLSQETFSDLWKLLPEN 18
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Result
No.
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Maximum DB
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OM protein - protein search, using sw model
    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                  Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    seq length: 0
seq length: 2000000000
    Query
Match Length
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47.832 Million cell updates/sec
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1: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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105
   GenCore version 5.1.9 Copyright (c) 1993 - 2006 Biocceleration Ltd.
   DВ
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 US-09-214-371-74
US-10-927-528-74
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US-110-155-059-11
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US-110-958-163R-3
US-110-918-643-3
US-110-918-643-3
US-110-918-63-31871
US-110-450-763-31871
US-110-450-763-31871
US-110-440-602-24
US-09-849-602-24
US-09-849-602-24
US-09-849-602-255R-8
US-110-466-255R-8
US-110-696-255R-8
US-10-696-255R-7
US-09-732-384-3
US-09-860-211-9
US-09-860-211-9
US-09-860-286-9
US-09-829-922-2
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US-10-274-874-4
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Sequence 74, Appl
Sequence 74, Appl
Sequence 8, Appli
Sequence 10, Appl
Sequence 16, Appl
Sequence 16, Appli
Sequence 3, Appli
Sequence 3, Appli
Sequence 33581, A
Sequence 31871, A
Sequence 24, Appl
Sequence 24, Appli
Sequence 7, Appli
Sequence 9, Appli
                                                                                                                                                                                                                                                                                                                Description
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RESULT 2
US-10-927-262A-74
; Sequence 74, Application US/10927262A
; Publication No. US20050137137A1
; GENERAL IMFORMATION:
APPLICANT: LANE, DAVID P
; APPLICANT: BOTTGER, VOLKER
APPLICANT: BOTTGER, ANGELIKA
APPLICANT: PICKSLEY, STEVEN M.
APPLICANT: HOCHKEPEL, HEINZ-KURT
APPLICANT: GARCIA-ECHEVERRIA, CARLOS
; APPLICANT: CHENE, PATRICK

B 6	US		
Query Match Best Local Matches 1	RESULT 1 Sequence 74, Application U Patent No. US20010018511A1 GENERAL INFORMATION: APPLICANT: Lane, David APPLICANT: Bottger, Volk APPLICANT: Bottger, Volk APPLICANT: Bottger, Volk APPLICANT: Chene, Patrid APPLICANT: Hochkeppel, H APPLICANT: Hochkeppel, H APPLICANT: Hochkeppel, H APPLICANT: Firet, Pascal TITLE OF INVENTION: Inhi FILE REFERENCE: 4-2037/ CURRENT APPLICATION NUMBER: APPLICANTON NUMBER: DATE: 199 PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1997 INUMBER OF SEQ ID NOS: 83 SOFTWARE: PATENTIN PATENTIN PATENTIN SEQ ID NO 74 LENGTH: 18 TYPE: PRT ORGANISM: Artificial Seq FEATURE: OTHER INFORMATION: Descr		2 2 2 2 2 2 2 2 2 2 2 2 3 2 3 3 3 3 3 3
	A-371-74 NO. USZOOLOOL B511A1 LINFORMATION: CANT: Lane, David CANT: Bottger, Volk CANT: Bottger, Volk CANT: Bottger, Volk CANT: Bottger, Ste CANT: Garcia-Echeve CANT: Garcia-Echeve CANT: Garcia-Echeve CANT: Garcia-Echeve CANT: Hochkeppel, H CANT: Hochkeppel, H CANT: Hochkeppel, H CANT: Garcia-Echeve CANT: Garcia-Echeve CANT: Garcia-Echeve CANT: Garcia-Echeve CANT: TULING DATE: 1937/ NT APPLICATION NUMBER: FILING DATE: 1997- APPLICATION NUMBER: FILING DATE: 1997- NO 74 TH: 18 INFORMATION: Descr WRS: PAT		
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larity Conserva SQETFSDL 	, Application US US20010018511A1 VORMATION: Lane, David Botteger, Volke Botteger, Angel Picksley, Step Chene, Patrick Hochkeppel, He Garcia-Echever Ferrete, Pascal INVENTION: Inhib ENCE: 4-20937/A PILICATION NUMBER: ING DATE: 1997-0 SEQ ID NOS: 83 PatentIN Ver. 2 8 Artificial Seque OCRMATION: Descri		ជាជាជាជាជាជាជាជាជាជាជាជាជាជាជាជាជាជា
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Score 99; DB 3; Len; Pred. No. 9.7e-08; O; Mismatches O; 18	14371B Kurt Carlos s of the Interaction of /09/214,371B 26 /EP97/03549 n of Artificial Sequence	ALIGNMENTS	US-10-160-290-2 US-10-077-176-54 US-10-077-176-55 US-10-077-176-56 US-10-077-176-57 US-10-434-693-2 US-10-165-216-4 US-10-165-216-4 US-10-191-121-3 US-10-391-068-2 US-10-391-068-2 US-10-391-071-9-20 US-10-31-31-31-31-31-31-31-31-31-31-31-31-31-
gth 18; Indels O; Gaps O	P53 and MDM2		Sequence 2, Appli Sequence 54, Appl Sequence 55, Appl Sequence 57, Appl Sequence 2, Appli Sequence 31, Appli Sequence 4, Appli Sequence 4, Appli Sequence 2, Appli Sequence 2, Appli Sequence 1, Appli Sequence 1, Appli Sequence 25, Appli Sequence 26, Appli Sequence 27, Appli Sequence 28, Appli Sequence 37, Appli Sequence 38, Appli Sequence 37, Appli Sequence 44, Appli

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RESULT 3
US-10-155-059-8
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Publication No. US20020147173A1
GENERAL INFORMATION:
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Best Local Similarity
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PRIOR FILING DATE: 1997-07-04
PRIOR APPLICATION NUMBER: GB 9614197.3
PRIOR FILING DATE: 1996-07-05
PRIOR APPLICATION NUMBER: GB 9707041.1
PRIOR FILING DATE: 1997-04-07
NUMBER OF SEQ ID NOS: 85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: INHIBITORS OF THE INTERACTION BETWEEN P53 AND MDM2 FILE REFERENCE: 39749.0002 APC CON CURRENT APPLICATION NUMBER: US/10/927,262A CURRENT FILING DATE: 2004-08-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 3.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH:
                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                 REGISTRATION NUMBER: 30,628
REFERENCE DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: METHODS OF TREATMENT USING NBS-1, ANTIBODIES AND PROTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                               APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 101 Federal Street
                                                                                                    NAME: Eisenstein, Ronald I
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FURET, PASCAL
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Pred. No.
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US-10-155-059-10
Sequence 10, Application US/10155059
Publication No. US20020147173A1
GENERAL INFORMATION
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                                                                                                                              US-10-155-059-10
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Best Local
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                                                                                                                                                                                                                                                      TELEFAX: 617-345-1300
TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
PILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
PILING DATE: 12-MAY-1997
ATTORNARY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FASESEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JOST, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 02110
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kaelin, William
Jost, Christine
                                                                                                                                              SEQUENCE DESCRIPTION: SEQ ID NO: 10:
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                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 30,628 REFERENCE/DOCKET NUMBER: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/10/155,059 FILING DATE: 24-May-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: MA
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                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                NAME: Eisenstein, Ronald I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
                                                                                                                                                                 TOPOLOGY: linear
                                                                                                                                                                                                                  LENGTH: 20 amino acids
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Conservative (
                                                        Conservative
                                                      94.3%; Score 99; DB 4; L
100.0%; Pred. No. 1.1e-07;
tive 0; Mismatches 0;
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100.0%; Pred. No. 1.1e-07;
Live 0; Mismatches 0;
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                                                                                            Length 20;
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RESULT 6
US-10-155-059-16
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US-10-155-059-11
                                                                                                                                                                                                      Sequence 16, Application US/101550
Publication No. US20020147173A1
GENERAL INFORMATION:
APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
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INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENCTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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GENERAL INFORMATION:
APPLICANT: Kaelin, William
Jost, Christine
Jost, METHOD!
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US/09/081,975
FILING DATE: 12-MAY-1998
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APPLICATION NUMBER: US/10/155,059
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                               JOST, Christine TITLE OF INVENTION: METHODS OF TREATMENT USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 11:
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MEDIUM TYPE: Diskette
                                          NUMBER OF SEQUENCES:
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CORRESPONDENCE ADDRESS:
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OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Eisenstein,
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                                                                                                                                                                                                                                                                                                                              Application US/10155059
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94.3%; Score 99; DB 4; 100.0%; Pred. No. 1.1e-
                                          ANTIBODIES : 27
                                                                                                                    NBS-1,
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                                                                                                 ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
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GENERAL INFORMATION:

APPLICANT: Hupp, Theodore
APPLICANT: Dornan, David
TITLE OF INVENTION: Screening Method and Agents
FILE REFERENCE: 9013.54
CURRENT APPLICATION NUMBER: US/10/467,758
CURRENT FILING DATE: 2003-08-13
PRIOR APPLICATION NUMBER: PCT/GB02/00640
PRIOR APPLICATION NUMBER: GB 0103508.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; SEQUENCE DESCRIPTION: SEQ ID NO: 16: US-10-155-059-16
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Best Local Similarity 100.0%; I
Marches 18; Conservative 0;
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SEQ ID NO 8
LENGTH: 20
TYPE: PRT
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INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 8, Application US/10467758 Publication No. US20040132108A1
                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 43
FEATURE:
NAME/KEY: MOD_RES
LOCATION: (5)...(5)
OTHER INFORMATION: RESIDUE MAY OPTIONALLY BE PHOSPHORYLATED
                                                                                               OTHER INFORMATION: EGFP BOX-1 PEPTIDE
                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <URNown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTMARE: FRACESCA for Windows Version 2.0b
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
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TELEFAX: 617-345-1300
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STREET: 101 Federal Street
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100.0%; Pred. No. 1.1
tive 0; Mismatches
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US-09-958-163A-3
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                                                                                                                                      Sequence 3, Application US/10918643 Publication No. US20050026143A1 GENERAL INFORMATION:
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Best Local (
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Best Local
FILE REFERENCE: sergeev CURRENT APPLICATION NUMBER: US/10/918,643 CURRENT FILING DATE: 2004-08-13 NUMBER OF SEQ ID NOS: 44 SOFTWARE: PatentIn version 3.2 SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: sergeev
CURRENT APPLICATION NUMBER: US/09/958,163A
CURRENT FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 44
                                                                                                     APPLICANT: Sergeev, Pavel TITLE OF INVENTION: Synthesis of biologically active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Sergeev, Pavel TITLE OF INVENTION: Synthesis of biologically active compounds in cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PUBLICATION INFORMATION:
AUTHORS: Harlow, E., Williamson, N.M., Ralston, R., Helfman, D.M. and AUTHORS: Adams, T.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: MOD_RES
LOCATION: (10)..(10)
OTHER INFORMATION: RESIDUE MAY OPTIONALLY BE PHOSPHORYLATED
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NAME/KEY: MOD_RES

LOCATION: (8)...(8)

OTHER INFORMATION: RESIDUE MAY OPTIONALLY BE PHOSPHORYLATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE: Molecular cloning and in vitro expression of a cDNA clone for TITLE: human cellular tumor antigen p53 JOURNAL: Molecular and cellular biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE: OTHER INFORMATION: part of the amino acid sequence of the Human tumour supressor p5
                                                                                                                                                                                                                                                                                                                                                                                                                              DATABASE ACCESSION NUMBER: K03199
DATABASE ENTRY DATE: 1995-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ISSUE: 7
PAGES: 1601-1610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VOLUME: 5
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Pred. No.
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Pred. No.
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APPLICANT: FOWLKES, Dana M.
APPLICANT: KAY, Brian K.
APPLICANT: KAY, Brian K.
APPLICANT: FRELINGER, Jeffrey A.
APPLICANT: FRELINGER, Jeffrey A.
APPLICANT: FRELINGER, Jeffrey A.
APPLICANT: FOR THE TOP TOWN THE TOWN THE TOWN THE TOWN OF DRUGS.
TITLE OF INVENTION: LIBRARIES
TITLE OF INVENTION: LIBRARIES
TITLE OF INVENTION: LIBRARIES
TITLE OF INVENTION: LIBRARIES
TITLE OF INVENTION: UNMEER: US/10/656,250
CURRENT FILING DATE: 1998-03-31
PRIOR APPLICATION NUMBER: 09/050,359
PRIOR FILING DATE: 1998-03-31
PRIOR APPLICATION NUMBER: 08/740,671
PRIOR APPLICATION NUMBER: 08/740,671
PRIOR APPLICATION NUMBER: 08/740,671
PRIOR FILING DATE: 1996-10-31
NUMBER OF SEQ ID NOS: 180
SOFTWARE: PATENTIN PATENTING 3.2
SEQ ID NO 48
LENGTH: 32
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                                                          Вb
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RESULT 11
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                                                                                                                                         Matches
                                                                                                                                                          Query Match
Best Local
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ISSUE: 7
PAGES: 1601-1610
DATE: 1985-07-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE: Molecular cloning and in vitro expression TITLE: human cellular tumor antigen p53 JOURNAL: Molecular and cellular biology VOLUME: 5
                                                                                                                                                                                                                                                                               ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                    TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DATABASE ACCESSION NUMBER: K03199 DATABASE ENTRY DATE: 1995-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                         OTHER INFORMATION: fragment of human p53
                                                                                                                                                                                                                                                            FEATURE:
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                                                      12 PPLSQETFSDLWKLLPEN 29
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                                                                               1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                            Similarity
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                                                                                                                                       Conservative
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100.0%; Pr/
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                                                                                                                                                            94.3%; Score 99;
100.0%; Pred. No.
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. 1.8e-07;
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US-10-450-763-33581

Sequence 33581, Application US/10450763 Publication No. US20050196754A1 GENERAL INFORMATION:

APPLICANT: Hyseq, Inc

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; LOCATION: (7)..(281)

OTHER INFORMATION: P53 domain identified by PFam, accession name P53, E-value=

OTHER INFORMATION: 1.4e-165, PFam score of 563.4

US-10-450-763-31871
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SOFTWARE: CUSTOM
SEQ ID NO 31871
LENGTH: 283
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Best Local :
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SOFTWARE: Custom
SEQ ID NO 33581
Query Match
Best Local Similarity
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TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
FILE REFERENCE: 790CIP3/US
CURRENT APPLICATION NUMBER: US/10/450,763
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: PCT/US01/08631
PRIOR FILING DATE: 2001-03-30
PRIOR FILING DATE: 2001-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
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PRIOR FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: PCT/US01/08631
PRIOR FILING DATE: 2001-03-30
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CURRENT FILING DATE: 2003-06-11
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                                                                                                                                                                                            LOCATION: (151)..(173)
OTHER INFORMATION: P53 TUMOUR SUPRESSOR SIGNATURE domain identified by eMATRIX,
OTHER INFORMATION: accession number PR00386B, p-value=5.950e-30, raw score of 12.28
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LOCATION: (97)..(137)
OTHER INFORMATION: p53 tumor antigen proteins domain identified by eMATRIX,
OTHER INFORMATION: accession number BL00348B, p-value=1.000e-40, raw score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (7)..(255)
OTHER INFORMATION: P53 domain identified by PFam,
OTHER INFORMATION: 5.1e-140, PFam score of 478.6
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                                                                                                                                                       NAME/KEY: DOMAIN
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                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
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LOCATION: (7)..(
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94.3%;
100.0%;
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Pred. No.
Score 99;
Pred. No.
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5. 1.6e-06;
0;
DB 5; I
. 1.8e-06;
                  Length 283
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Publication No. US20030165834A1
GENERAL INFORMATION:
APPLICANT: Scanlan, Matthew J.
APPLICANT: Old, Lloyd J.
APPLICANT: Stockert, Elisabeth
APPLICANT: Chen, Yao-Tseng
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                                                                                                                                                                                               SOFTWARE: PatentIn version 3.0
SEQ ID NO 24
SENGTH: 353
TYPE: PRT
ORGANISM: Homo sapiens
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APPLICANT: Deppert, Wolfgang W.
APPLICANT: Dornreiter, Irene
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Publication No. US20060062792A1
                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/11/236,014
CURRENT FILING DATE: 2005-09-27
PRIOR APPLICATION NUMBER: PCT/EP2004/003299
PRIOR FILING DATE: 2004-03-27
PRIOR APPLICATION NUMBER: EP 03007000.7
PRIOR PILING DATE: 2003-03-27
PRIOR FILING DATE: 2003-03-27
NUMBER OF SEQ ID NOS: 26
                                                                                                                                     FEATURE:
NAME/KEY: UNSURE
LOCATION: (76)...(76)
OTHER INFORMATION: 1
                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Colon Cancer Antigen | FILE REFERENCE: L0461/7105 (JRV) CURRENT APPLICATION NUMBER: US/09/849,602 CURRENT FILING DATE: 2001-05-04
                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Novel Human p53 Splice Variant Displaying Differential TITLE OF INVENTION: Transcriptional Activity FILE REFERENCE: 4121-178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn version 3.3
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TYPE: PRT
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                             1 PPLSQETFSDLWKLLPEN 18
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                                                                            Similarity
PPLSQETFSDLWKLLPEN
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To. US20030165834A1
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                                                            Conservative
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                                                                                                                                     X = any amino acid
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100.0%; Pr
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                                                                            Score 99;
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Pred. No.
                                                            Mismatches
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                                                                            DB 3; L
2.3e-06;
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US-10-146-473-78
US-10-146-473-78
US-10-146-473-78
Sequence 78, Application US/10146473
PULICANT NORMATION:
APPLICANT Gout, Ivan
APPLICANT Gout, Ali
APPLICANT Gou
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 s
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OM protein -
        8
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seq
                                               protein search, using sw model
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length: 2000000000
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Gapop 10.0 ,
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105
      Published Applications AA New:*

| MemC_Celerra_SIDS3/prodata/1/pubpaa/US09_NEW_PUB.pep:*
| EMC_Celerra_SIDS3/prodata/1/pubpaa/US06_NEW_PUB.pep:*
| MemC_Celerra_SIDS3/prodata/1/pubpaa/US07_NEW_PUB.pep:*
| MemC_Celerra_SIDS3/prodata/1/pubpaa/US08_NEW_PUB.pep:*
| MemC_Celerra_SIDS3/prodata/1/pubpaa/US08_NEW_PUB.pep:*
| MemC_Celerra_SIDS3/prodata/1/pubpaa/US11_NEW_PUB.pep:*
| MemC_Celerra_SIDS3/prodata/1/pubpaa/US11_NEW_PUB.pep:*
| MemC_Celerra_SIDS3/prodata/1/pubpaa/US11_NEW_PUB.pep:*
| MemC_Celerra_SIDS3/prodata/1/pubpaa/US11_NEW_PUB.pep:*
| MemC_Celerra_SIDS3/prodata/1/pubpaa/US10_NEW_PUB.pep:*
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      GenCore version 5.1.9 (c) 1993 - 2006 Biocceleration Ltd.
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                       US-10-538-066-367

US-11-315-777-9

US-11-319-873-9

US-11-319-873-9

US-11-009-357-6

US-11-09-357-6

US-10-953-613C-79

US-10-953-613C-78

US-10-953-613C-78

US-10-949-953-613C-78

US-10-949-953-613C-78

US-10-953-613C-78

US-10-949-953-613C-78

US-10-949-953-613C-78

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US-10-949-953-613C-78

US-10-949-953-613C-78

US-10-949-953-633

US-10-449-902-5579

US-09-949-955-241

US-10-449-902-5579

US-10-449-902-5579

US-10-449-902-5579

US-10-449-902-5579

US-11-293-697-3673
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Sequence 367, Applisequence 3, Applisequence 3, Applisequence 2, Applisequence 778, Applisequence 778, Applisequence 780, Applisequence 780, Applisequence 780, Applisequence 791, Applisequence 791, Applisequence 624, Applisequence 624, Applisequence 610, Applisequence 781, Applisequence 50286, Asequence 50286, Asequence 241, Applisequence 241, Applisequence 241, Applisequence 241, Applisequence 3028, Applisequence 3028, Applisequence 313, Applisequence 313, Applisequence 313, Applisequence 313, Applisequence 3673, Applis
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US-11-315-777-9
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US-10-538-066-367
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Best Local S
Matches 18
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42.9 38.6 US-10-953-349-5776 44.9 38.6 US-10-953-349-5775 44.41.9 43.9 6 US-10-449-902-33353 44.41.9 72.6 6 US-10-449-902-5651.5 44.41.9 72.6 6 US-10-449-902-5651.5 44.41.9 72.6 6 US-10-449-902-5651.5 44.41.9 72.6 6 US-10-449-902-5650 43.41.0 10.0 10.8-10-449-902-49787 42.5 40.0 11.3 7 US-11-293-697-2548 42.40.0 81.6 US-10-953-349-18620 42.40.0 81.6 US-10-953-349-18620 42.40.0 113.6 US-10-953-349-18617 42.40.0 24.4 6 US-10-953-349-18617 42.40.0 25.4 6 US-10-953-349-18617 42.40.0 25.4 6 US-10-953-349-18617 42.40.0 25.4 6 US-10-953-349-18617 42.40.0 25.6 US-10-953-349-20974 42.40.0 25.6 US-10-953-349-20975 42.40.0 25.6 US-10-953-349-20975 42.40.0 25.6 US-10-953-349-35956 42.40.0 53.8 6 US-10-953-349-35955	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27
381 6 US-11 381 6 US-12 381 6 US-12 39 439 6 US-12 39 726 6 US-12 39 726 6 US-12 372 6 US-12 373 6 US-12 374 6 US-12 375 6 US-12	42	42	42	42	42	42	42	42	42	42.5	43	43	44	44	44	44	44	45	45
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	US-10-953-349-35955	-10-953	1	-10	Ξ	-10-953	-10-953	-10-953	-10	-11	Ξ	ï	Ξ	L	L	-	-10	Ľ	US-10-953-349-5776
	35955, A	35956, A	20973, A	46136, A	20974, A	20975, A	18617, A	18620, A	354, App	2548, Ap	49787, A	623, App	56500, A	56115, A	51533, A	42702, A	33353, A	5775, Ap	5776, Ap

ALIGNMENTS

GENERAL INFORMATION: APPLICANT: Epimmune Inc. TITLE OF INVENTION: Peptides and Compositions TITLE OF INVENTION: Peptides and Compositions FILE REFERENCE: 2060.015PC06 CURRENT APPLICATION UNMBER: US/10/538,066 CURRENT FILING DATE: 2005-06-09 PRIOR APPLICATION NUMBER: US 60/432,017 PRIOR TILING DATE: 2002-12-10 NUMBER OF SEQ ID NOS: 767 SOFTWARE: PatentIn version 3.2 SEQ ID NO 367 LENGTH: 393 TYPE: PRI OPCANION. ; ORGANISM: Homo sapiens US-10-538-066-367 Sequence 9, Application US/11315777 Publication No. US20060099187A1 GENERAL INFORMATION: APPLICANT: Gregory, Richard J. Wills, Ken N. Sequence 367, Application US/10538066 Publication No. US20060094649A1 NUMBER OF SEQUENCES: 9 CORRESPONDENCE ADDRESSE: Townsend and STREET: Two Embarcadero CITY: San Francisco Maneval, Daniel C. TITLE OF INVENTION: Recombinant Adenoviral Vector and Methods of Use 1 PPLSQETFSDLWKLLPEN 18 Similarity STATE: California COUNTRY: USA ZIP: 94111-3834 PPLSQETFSDLWKLLPEN Conservative 94.3%; 50 100.0%; Pr 29 Score 99; DB 6; Length 393; pred. No. 2.9e-07; Mismatches Townsend and Crew LLP Center, Eighth Floor -B7, and -B44 Tumor 0, Associated Antigen Gaps 0

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; TYPE: PRT; ORGANISM: Homo sapien US-11-340-715-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/11340715 Publication No. US20060115852A1 GENERAL INFORMATION:
                                                                                      NUMBER OF SEQ ID NOS: 29
SOFTWARE: PatentIn version 3.2
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                            APPLICANT: IKAWA, Yoji
APPLICANT: IKAWA, Shuntaro
APPLICANT: OBINATA, Masuo
TITLE OF INVENTION: HUMAN P51 GENES AND GENE PRODUCTS THEREOF
FILE REFERENCE: Q61014
                                                                                                                                                     PRIOR APPLICATION NUMBER: JP 10-100467
PRIOR FILING DATE: 1998-03-27
PRIOR APPLICATION NUMBER: PCT/JP99/01512
PRIOR FILING DATE: 1999-03-24
                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/11/340,715
CURRENT FILING DATE: 2006-01-27
PRIOR APPLICATION NUMBER: US/09/670,568
PRIOR FILING DATE: 2001-01-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
                                                                 LENGTH: 393
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FILING DATE: 18-May-2001
APPLICATION NUMBER: US/08/328,673
FILING DATE: 25-OCT-1994
APPLICATION NUMBER: US 08/142,669
FILING DATE: 25-OCT-1993
APPLICATION NUMBER: US 08/233,669
FILING DATE: 26-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: 1..393
OTHER INFORMATION: /note= "human p53"
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/315,777
FILING DATE: 21-Dec-2005
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Smith, Timothy S.
REGISTRATION NUMBER: 35,367
REFERENCE/DOCKET NUMBER: 016930-000920US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12 PPLSQETFSDLWKLLPEN 29
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 393 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94.3%; Score 99; DB 7; I
100.0%; Pred. No. 2.9e-07;
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US-11-009-357-6
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 6
LENGTH: 393
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Publication No. US20060127376A1
GENERAL INFORMATION:
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Publication No. US20060140910A1
GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
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CURRENT FILING DATE: 2004-12-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Moll, Ute
TITLE OF INVENTION: Methods and Compositions for Modulating Apoptotic Pathways
FILE REFERENCE: STONYB-09615
            COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/11/319,873
FILING DATE: 27-Dec-2005
CLASSIFICATION LOTA:
APPLICATION NUMBER: US/11/319,873
FILING DATE: 27-Dec-2005
PILING DATE: 27-Dec-2005
FILING DATE: 28-Oct-1997
APPLICATION NUMBER: US/08/958,570
APPLICATION NUMBER: US/08/328,673
FILING DATE: 25-Oct-1994
APPLICATION NUMBER: US/08/328,673
APPLICATION NUMBER: US/08/328,669
FILING DATE: 25-Oct-1993
APPLICATION NUMBER: US/08/233,669
                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Recombinant Adenoviral Vector Methods of Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gregory, Richard J. Wills, Ken N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12 PPLSQETFSDLWKLLPEN 29
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                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                     STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: San Francisco
FILING DATE: 26-APR-1994
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100.0%; Pred. No.
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2.9e-07;
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Publication No. US20060127376A1
GENERAL INFORMATION:
APPLICANT: Moll, Ute
TITLE OF INVENTION: Methods and Compositions for Modulating Apoptotic Pathways
FILE REFERENCE: STONYS-05615
CURRENT APPLICATION NUMBER: US/11/009,357
CURRENT FILING DATE: 2004-12-10
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin version 3.3
SEQ ID NO 2
Sequence 778, Application US/10953613C
Publication No. US20060127404A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon;
APPLICANT: Bernard, Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CEN5038 NP
CURRENT FILLING DATE: US/10/953,613C
CURRENT FILLING DATE: 2004-09-29
PRIOR APPLICATION NUMBER: 60/507,231
PRIOR APPLICATION NUMBER: 60/507,231
PRIOR FILING DATE: 2003-09-30
NUMBER OF SEQ ID NOS: 1021
SOFTWARE: Patentin Ver 3.0
SEQ ID NO 778
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Best Local :
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Smith, Timothy S.
REGISTRATION NUMBER: 35,367
REFERENCE/DOCKET NUMBER: 016930-000920US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
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LOCATION:
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TOPOLOGY: linear
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100.0%; Pr
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0. 2.9e-07;
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; ORGANISM: Homo sapiens
US-10-953-613C-780
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US-10-953-613C-790
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                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/953,613C CURRENT FILING DATE: 2004-09-29 PRIOR APPLICATION NUMBER: 60/507,231 PRIOR FILING DATE: 2003-09-30 NUMBER OF SEQ ID NOS: 1021 SOFTWARE: Patentin Ver 3.0 SEQ ID NO 780
                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon;
APPLICANT: Bernard; Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CENSO38 NP
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CURRENT APPLICATION NUMBER: US/10/953,613C
CURRENT FILING DATE: 2004-09-29
PRIOR APPLICATION NUMBER: 60/507,231
PRIOR FILING DATE: 2003-09-30
NUMBER OF SEQ ID NOS: 1021
SOFTWARE: PATENTIN Ver 3.0
SEQ ID NO 790
                                  Matches
                                                   Query Match
Best Local (
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Best Local Similarity
Matches 12; Conserv
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TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
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5 OETFSDLWKLLP 16
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                                                   Similarity
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                                  Conservative
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91.7%;
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100.0%; Pred. No. 0.00056;
tive 0; Mismatches 0;
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                                                 Score 61; DB 6;
Pred. No. 0.003;
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Pred. No.
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QETFSDYWKLLP 12

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RESULT 12
US-10-933-613C-791
y Sequence 791, Application US/10953613C
Publication No. US20060127404A1
GENERAL INFORMATION:
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US-10-953-613C-792
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US-10-953-613C-779
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Publication No. US20060127404A1
GENERAL INFORMATION:
APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon;
APPLICANT: Bernard; Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CEN5038 NP
FULE REFERENCE: COMPOSITION NUMBER: US/10/953,613C
CURRENT FILING DATE: 2004-09-29
PRIOR FILING DATE: 2003-09-30
PRIOR FILING DATE: 2003-09-30
PRIOR FILING DATE: 2003-09-30
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Best Local Similarity
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               APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, APPLICANT: Bernard; Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHO
FILE REFERENCE: CEN5038 NP
CURRENT APPLICATION NUMBER: US/10/953,613C
CURRENT FILING DATE: 2004-09-29
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TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CEN5038 NP
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PRIOR APPLICATION NUMBER: 60/507,231
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91.7%;
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91.7%;
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Pred. No. 0.0042;
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Pred. No. 0.003;
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                                                                                                                          John;Scallon;
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RESULT 14
US-10-538-066-600
j Sequence 600, Application
j Publication No. US20060094
j GENERAL INFORMATION:
j APPLICANT: Epimmune Inc.
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                                                                             ; ORGANISM: Homo sapiers US-10-538-066-600
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                                                                                                                               TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and -B44 Tumor Associated Antigen TITLE OF INVENTION: Peptides and Compositions FILE REFERENCE: 2060.015PC06 CURRENT APPLICATION NUMBER: US/10/538,066 CURRENT FILING DATE: 2005-06-09 PRIOR APPLICATION NUMBER: US 60/432,017 PRIOR APPLICATION NUMBER: US 60/432,017 PRIOR FILING DATE: 2002-12-10 NUMBER: US 60/432,017 PRIOR FILING DATE: 2002-12-10 SEQ ID NO 600 LENGTH: 10 CONTRACT OF THE PRIOR APPLICATION VERSION 3.2
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SEQ ID NO 624
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Publication No. US20060094649A1
GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 1021
SOFTWARE: PatentIn Ver 3.0
SEQ ID NO 791
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APPLICANT: Epimmune Inc.

TITLE OF INVENTION: Publishes and Compositions

TITLE OF INVENTION: Peptides and Compositions

FILE REFERENCE: 2060.015PC06

CURRENT APPLICATION NUMBER: US/10/538,066

CURRENT EPILING DATE: 2005-06-09

PRIOR APPLICATION NUMBER: US 60/432,017

PRIOR PILING DATE: 2002-12-10

NUMBER OF SEQ ID NOS: 767

NUMBER OF SEQ ID NOS: 767
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Best Local Similarity
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TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                   TYPE: PRT
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  Conservative
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52.4%; Score 55; DB 6;
100.0%; Pred. No. 0.019;
tive 0; Mismatches
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91.7%;
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Pred. No. 0.0042;
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RESULT 15

US-10-953-613C-781

US-10-953-613C-781

Sequence 781, Application US/10953613C

Publication No. US20060127404A1

GENERAL INFORMATION:

APPLICANT: Huang, Chichi, Heavner, George; Knight, David; Ghrayeb, John; Scallon;

APPLICANT: Bernard; Nesspor; Thomas

TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES

FILE REFERENCE: CENSO38 NP

CURRENT APPLICATION NUMBER: US/10/953,613C

CURRENT APPLICATION NUMBER: 60/507,231

PRIOR APPLICATION NUMBER: 60/507,231

PRIOR FILING DATE: 2003-09-30

NUMBER OF SEQ ID NOS: 1021

SOFTWARE: Patentin Ver 3.0

SEQ ID NO 781

LENGTH: 12

TYPE: PRT

ORGANISM: Homo sapiens

US-10-953-613C-781
Search completed: July 5, 2006, 22:51:02 Job time : 20 secs
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                                                                                                                                                                           Query Match 52.4%; Score 55; DB 6; Length 12; Best Local Similarity 83.3%; Pred. No. 0.023; Matches 10; Conservative 0; Mismatches 2; Indels
                                                                                                              5 QETFSDLWKLLP 16
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Maximum DB
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Perfect score:
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Match Length
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| / EMC Celerra SIDS3/ptodata/2/iaa/5 COMB.pep:*

| EMC Celerra SIDS3/ptodata/2/iaa/6 COMB.pep:*

| EMC Celerra SIDS3/ptodata/2/iaa/7 COMB.pep:*

| / EMC Celerra SIDS3/ptodata/2/iaa/H COMB.pep:*

| EMC Celerra SIDS3/ptodata/2/iaa/FECOMB.pep:*

| EMC Celerra SIDS3/ptodata/2/iaa/RE COMB.pep:*

| EMC Celerra SIDS3/ptodata/2/iaa/RE COMB.pep:*

| EMC Celerra SIDS3/ptodata/2/iaa/BECOMB.pep:*
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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US-09-08-975-48
US-09-958-163A-3
US-08-390-546-1
US-08-390-516C-1
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US-08-897-221-19
US-09-603-052-2
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US-08-697-221-19
US-08-697-221-20
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Sequence 11, Appli
Sequence 3, Appli
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Sequence 1, Appli
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94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3
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Query Match 94.3
Best Local Similarity 100
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FESSESQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 60/046,207
                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                             NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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CITY: Boston
STATE: MA
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ZIP: 021
                                                                                                                              TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                              TOPOLOGY:
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94.3%; Score 99; DB 2; L
100.0%; Pred. No. 4.3e-09;
/ative 0; Mismatches 0;
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RESULT 3
US-09-081-975-11
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US-09-081-975-10
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Best Local S
Matches 18
Sequence 11, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTBINS THERETO, AND USES
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTBINS THERETO, AND USES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617,345-6654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: UZLIV
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FASTSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
CONTINUATION NUMBER: US/09/081,975
                                                                                                                                                                                                                                                                                                                                                                                                            LD NO:
LIPE: amino acid
STRANDEDNESS: si-
TOPOLOGY: 1'
081-975-
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COUNTRY: US
ZIP: 02110
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CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
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6451979
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100.0%; Pred. No. 4.3e-09;
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RESULT 4
US-09-081-975-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Versic
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CILASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
APPLICATION NUMBER: 60/046,207
                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTI
TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compartible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STAIL COUNTRY: COUNTRY: 02110
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nes 18; Conservative (
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REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 4:
                                                                                                                                                                                                                                     ZIP:
                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                         CITY: Boston
                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 101 | CITY: Boston
                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Nixon Peabody LLP
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100.0%; Pred. No. 4.3e-09;
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RESULT 5
US-09-069-827A-48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 48, Application US/09069827A Patent No. 6617114 GENERAL INFORMATION:
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                                                    INFORMATION FOR SEQ ID NO: 48: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 617-345-6054
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                                                                                                                                                                                                                                                                                                                                  COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/069,827A
FILING DATE: 30-Apr-1998
CLASSIFICATION: cUnknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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STRANDEDNESS: sir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Eisenstein, Ro
REGISTRATION NUMBER:
                                                                                                      NAME: COOPER, Iver P
REGISTRATION NUMBER: 28,005
REFERENCE/DOCKET NUMBER: FOWLKES=4C
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
                                                                                                                                                                                              APPLICATION NUMBER: US 08/740,671 FILING DATE: 31-OCT-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 178 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FRELINGER, Jeffrey A
HYDE-DERUYSCHER, RODIN P
TITLE OF INVENTION: IDENTIFICATION OF DRUGS USING
COMPLEMENTARY COMBINATORIAL LIBRARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: FOWLKES, Dana M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
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                                                                                                                                                                                                                                                APPLICATION NUMBER: US 09/050,359
FILING DATE: 31-MAR-1998
APPLICATION NUMBER: PCT/U897/19638
FILING DATE: 31-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C. STREET: 624 Ninth Street N.W., Suite 300
                  LENGTH: 32 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: D.C.
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PPLSQETFSDLWKLLPEN 19
                                                                      TELEPHONE: (202) 737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 amino acids
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RESULT 7
US-08-245-500A-1
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; SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-069-827A-48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-958-163A-3
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                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: BURRELL
APPLICANT: HILL, D
APPLICANT: KINCLER
APPLICANT: VOGELST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin version 3.2
SEQ ID NO 3
LENGTH: 32
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Patent No. 6831071
GENERAL INFORMATION:
APPLICANT: Sergeev, Pavel
TITLE OF INVENTION: Synthesis of biologically active compounds in cells
FILE REFERENCE: sergeev
                                                                                                                                                                                                                                                         Sequence 1, Application US/08245500A Patent No. 5550023
                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PEATURE:
OTHER INFORMATION: part of the amino acid sequence of the Human tumour supressor p5
PUBLICATION INFORMATION:
AUTHORS: Harlow,E., Williamson,N.M., Ralston,R., Helfman,D.M. and
AUTHORS: Adams,T.E.
TITLE: Molecular cloning and in vitro expression of a cDNA clone for
TITLE: human cellular tumor antigen p53
TITLE: human cellular tumor antigen p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/958,163A CURRENT FILING DATE: 2001-10-03
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TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DATE: 1985-07-01
DATABASE ACCESSION NUMBER: K03199
DATABASE ENTRY DATE: 1995-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ISSUE: 7
PAGES: 1601-1610
                                                                                                               TITLE OF INVENTION: AMPLIFICATION TITLE OF INVENTION: HUMAN TUMORS NUMBER OF SEQUENCES: 5
                                                                                                  CORRESPONDENCE ADDRESS:
              STATE: D.C.
COUNTRY: USA
                                               STREET: 1001 G STREET: WASHINGTON
                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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18; Conserv
20001
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                                                                 1001 G STREET, N.W.
                                                                                                                                                                      KINZLER, KENNETH
VOGELSTEIN, BERT
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                                                                                                                                                                                                                          BURRELL, MARILEE
                                                                                  BANNER, BIRCH, MCKIE AND BECKETT
                                                                                                                                                                                                        DAVID E.
                                                                                                                                                                                      KENNETH W.
                                                                                                                                                       AMPLIFICATION OF HUMAN MDM2 GENE IN
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100.0%; Pred. No.
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%; Pred. No. 7.1
0; Mismatches
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US-08-390-546-1
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                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: BURRELL
APPLICANT: HILL, D
APPLICANT: KINZLER
APPLICANT: VOGELST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IELEX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 64 amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local Similarity
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ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
                                                                                                                                                                                                                                                                                                                                    APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
TITLE OF INVENTION: HUMAN TUMORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: US,
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                             APPLICATION NUMBER: US/08/390,546
FILING DATE: 07-APR-1993
CLASSIFICATION: 536
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                    CITY: WASHINGTON
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CLASSIFICATION: 435
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                                                                                                                                                                                                                                   D.C.
                                                                                                                                                                                                                                                                 E: BANNER, BIRCH, MCKIE AND BECKETT 1001 G STREET, N.W.
                                                                                                                                                                                                                     USA
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NO
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100.0%; Pred. No.
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Patent No. 5618921
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HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 197430 BBMB UT INFORMATION FOR SEQ ID NO:
                                                                              TELEX: 197430 BBMB UT INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,479A
                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: HILL, DAVID B.
APPLICANT: KINGLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
APPLICANT: VOGELSTEIN, BERT
APPLICANT: TAVENTION: AMPLIFICATION OF HUMAN MDM2 GENE
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ADDRESSEE: BANNER & WITCOFF, LTD
                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 1001 G ST
CITY: WASHINGTON
STATE: D.C.
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                                                                                                                                                                                                                                                     FILING DATE: 02
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 64 amino acids
TYPE: amino acid
STRANDEDNESS: single
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TELEFAX: 202-508-9299
                                              LENGTH: 64 amino acids
TYPE: amino acid
                                                                                                                               TELEPHONE: 202-508-91
TELEFAX: 202-508-9299
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               TOPOLOGY: lir
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protein
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; Pred. No.
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 64 amin
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Best Local :
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APPLICATION NUMBER: US 08/245,500
FILING DATE: 18-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.4279.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
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APPLICANT: BURREI
                                                                                            FRAGMENT TYPE: N
ORIGINAL SOURCE:
ORGANISM: Home
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                               MOLECULE NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens POSITION IN GENOME: CHROMOSOME/SEGMENT: 17q
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                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
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TELEPHONE: Z
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                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
TOPOLOGY: lir
CHROMOSOME/SEGMENT: 17q
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amino acid
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linear
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                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01107.42798
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                                                      Matches
                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 011
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION: APPLICANT: BURREL
                                                                                                                                                                                                                                                                                                            TELEX: 197430 BBMB UT INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                           TOPULOGA: p:
MOLECULE TYPE: p:
HYPOTHETICAL: NO
                                                                                                                                      POSITION IN GENOME:
                                                                                                                                                                  FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 07-APCLASSIFICATION: 53
                                                                                                                                                                                                                                                   LENGTH: 64 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                     TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                        CHROMOSOME/SEGMENT: 17q
                                                                                                                                                      ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
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12 PPLSQETFSDLWKLLPEN 29
                                                      18;
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                    1 PPLSQETFSDLWKLLPEN 18
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                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KINZLER, KENNETH W. VOGELSTEIN, BERT
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                                                                                                                                                                               N-terminal
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                                               94.3%; but
100.0%; Pr
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100.0%; Pr
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                                                                  Score 99; DB 1; I
; Pred. No. 1.5e-08;
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RESULT 12 US-08-390-517A-1

; Sequence 1, Application US/08390517A

Patent No. 5736338

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                                                                                                                                                                                            Sequence 1, Application US/08390515A Patent No. 5756455
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Best Local
                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: BURREL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 197430 BBMB UT INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens POSITION IN GENOME: CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                   APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: BANNER, BIRCH, M
STREET: 1001 G STREET, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 202-508-9100 TELEFAX: 202-508-9299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: KINZLER, KENNETH W. APPLICANT: VOGELSTEIN, BERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: BANNER, BIRCH, MCKIE AND BECKETT STREET: 1001 G STREET, N.W.
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 07-APR CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 64 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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                                                                                                                                                                                                                                                                                                          12 PPLSQETFSDLWKLLPEN 29
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07-APR-1993
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                BIRCH, MCKIE AND BECKETT
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thes 0;
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; Patent No. 5858976
; GENERAL INFORMATION:
APPLICANT: BURRELL
; APPLICANT: HILL, D
; APPLICANT: KINZLER
; APPLICANT: KINZLER
                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-801-718-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 14
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           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Ve:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/801,718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 197430 BBMB UT INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 64 amino acida
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,515A
FILING DATE: 07-APR-1993
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FRAGMENT TYPE: N-terminal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                           TITLE OF INVENTION: AMPLIFICATION TITLE OF INVENTION: HUMAN TUMORS NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: NO ANTI-SENSE: NO
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COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 64 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                      COUNTRY: USA
ZIP: 20001
                                                                                                                                                                          STATE: D.C
                                                                                                                                                                                                             STREET:
                                                                                                                                                                                            CITY:
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FILING DATE:
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                                                                                                                                                                                                                                                                                                                                   HILL, DAVID E.
KINZLER, KENNETH W.
                                                                                                                                                                                                                                                                                                                    VOGELSTEIN, BERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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14-FEB-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94.3%; Score 99; DB 1;
100.0%; Pred. No. 1.5e-
cive 0; Mismatches
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                                                                                                                                                                                                                                                                                                  OF HUMAN MDM2 GENE
                                                   Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.5e-08;
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RESULT 15
US-09-170-159A-1
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 0110
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
TELEPAX: 202-508-9299
TELEPAX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 64 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
POSITION IN GENOME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/09170159A Patent No. 6399755
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/390,515
PILING DATE: 07-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/170,159A
FILING DATE: 13-Oct-1998
CLASSIFICATION: <UNknown>
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42798
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: BANNER, BIRCH, MCKIE AND BECKETT STREET; 1001 G STREET, N.W.
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BURRELL, MARILEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity 100.
18; Conservative
TELEPHONE: 202-508-9100
TELEFAX: 202-508-9299
TELEX: 197430 BBMB UT
                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 20001
                                                                                                                                                                                                                                                                                                                                                                     STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KINZLER, KENNETH W. VOGELSTEIN, BERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94.3%; Scc.
100.0%; Pr
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; Pred. No. 1.5e-0
0; Mismatches
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                                                                    Matches 18; Conservative
                                                                                                   Query Match
                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                  POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                          HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: protein
 12 PPLSQETFSDLWKLLPEN
                   1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                             LENGTH: 64 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                         ORGANISM: Homo sapiens
                                                                  94.3%; Score 99; DB 2; Length 64; 100.0%; Pred. No. 1.5e-08; tive 0; Mismatches 0; Indels
29
                                                                    0;
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Gaps

0,

Search completed: July 5, 2006, 22:47:26 Job time : 51 secs

0,

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Database : P 1: 2: 3: 4:	Post-processing: Minimum Match Maximum Match Listing first	Minimum DB seg length: 0 Maximum DB seg length: 19	Total number of hi	Searched: 28	Scoring table: BL Ga	Title: US- Perfect score: 105 Sequence: 1 P	Run on: Ju	OM protein - prote	
PIR 80:* : pir1:* : pir2:* : pir3:* : pir4:*	Minimum Match 0% Maximum Match 100% Listing first 45 summaries	gth: 0 gth: 19	Total number of hits satisfying chosen parameters: 3436	283416 segs, 96216763 residues	BLOSUM62 Gapop 10.0 , Gapext 0.5	US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19	July 5, 2006, 22:54:41; Search time 38 Seconds (without alignments) 48.108 Million cell updates/sec	OM protein - protein search, using sw model	GenCore version 5.1.9 Copyright (c) 1993 - 2006 Biocceleration Ltd.

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	ø	8	7	o,	5	4.	w	N	1	Result No.
21	21	21	21	21	21	21	21	21.5	21.5			22	22	22	22	22	22	22	22	23	23	23	24	24		25			Score
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beta-D-galactosida	214K exoantigen (v	neuropeptide Pec-H	hypertrehalosemic	hypertrehalosemic	adipokinetic hormo	hypertrehalosemic	μ.		T cell receptor al	Ig H chain V-D-J r		insulin-like growt	obuty	lymnaDFamide 5 - g	4	2		hypothetical prote	×	protein kinase (EC		adipokinetic hormo	chaperonin 10 homo		translation elonga	etica.	\mathbf{r}	ome polyprote	Description

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19.0	19.0	19.0	19.0	19.5	19.5	19.5	20.0	20.0	20.0	20.0	20.0	20.0	20.0	20.0	20.0
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A49823	S08995	A44960	A33995	I52623	PH1627	PC4391	I49037	A35704	S15754	E58503	A35389	A45103	B56864	B35245	A35245
adipokinetic hormo	hypertrehalosemic	neuropeptide Led-C	adipokinetic hormo	hypothetical prote	Ig H chain V-D-J r	cysteine proteinas	TcR delta chain V-	cytochrome P450 ol	actin 6 - soybean	superoxide dismuta	urease (EC 3.5.1.5	7 alpha-hydroxy-4-	dipeptidyl-peptida	histone H1.c - mou	histone Hla - mous

ALIGNMENTS

genome polyprotein (clone L3/S2) - Skalica virus (fragment)
C;Species: Skalica virus
C;Species: Skalica virus
C;Accession: B61409
R;Giirakhoo, F.; Heinz, F.X.; Mandl, C.W.; Holzmann, H.; Kunz, C.; Gresikova, M.
J. Gen. Virol. 72, 333-338, 1991
A;Title: The relationship between the flaviviruses Skalica and Langat as revealed by mo A;Reference number: A61409; MUID:91132129; PMID:1847173
A;Accession: B61409
A;Status: preliminary; not compared with conceptual translation
A;Status: preliminary; not compared with conceptual translation
A;Residues: 1-19 -GUI's
A;Cross-references: UNIPROT:Q7LZY3; UNIPARC:UPI0000174A0E
C;Superfamily: hepatitis C virus genome polyprotein

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Tha p 1 - Thaumetopoea pityocampa (fragment)
C;Species: Thaumetopoea pityocampa
C;Date: 03-Jun-2002 #sequence_revision 03-Jun-2002 #text_change 09-Jul-2004
C;Accession: A59396
R;Moneo, I.
submitted to the Protein Sequence Database, September 2001
A;Description: Isolation and characterization of a major allergen of the pine procession, Reference number: A59396
A;Accession: A59396
A;Accession: A59396
                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 2
A59396
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                                                                                                                                A;Cross-references: UNIPROT:Q7M4K8; UNIPARC:UPI000017CD37
A;Experimental source: L5
A;Note: IgB-binding protein, major allergen
                                                                                                                                                                                     A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-18 <MON>
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Best Local Similarity 41.7%;
Matches 5; Conservative
                                                                                  Query Match
Best Local Similarity
                                                                     Matches
2 ETYSDKYDTIDVN 14
                                  6 ETFSDLWKLLPEN 18
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Pred. No. 1.3e+03;
3; Mismatches 5;
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Pred. No. 84;
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R;Saville, B.J.; Collins, R.A.
Cell 61, 685-696, 1990
A;Title: A site-specific self-cleavage reaction performed A;Reference number: A35105; MUID:90263093; PMID:2160856
A;Accession: A35105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: mitochondrion Neurospora crassa mitochondrion (rragment)
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 07-Dec-1999
C;Accession: A35105
R;Saville, B.J.; Collins, R.A.
                                                                                   Biochem. Biophys. Res. Commun. 155, 344-350, 1988
A;Title: Isolation and primary structure of a neuropeptide
A;Reference number: A31571; MUID:88326324; PMID:3415690
A;Accession: A31571
A;Molecule type: protein
A;Residues: 1-10 <JAF>
A;Cross_references: UNIPROT:P16353; UNIPARC:UPI000012CDC5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Kamo, M.; Kawakami, T.; submitted to JIPID, March A;Reference number: PA0109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          translation elongation factor eEF-1 beta' chain - Arabidopsis thaliana (fragment) (;Species: Arabidopsis thaliana (mouse-ear cress) (;Decies: Arabidopsis thaliana (mouse-ear cress) (;Date: 07-Apr-1995 #sequence_revision 26-May-1995 #text_change 31-Dec-2004 (;Accession: PA0110
A;Residues: i=10 CJAF>
A;Cross-references: UNIPROT:P16353; UNIPARC:UPI000012CDC5
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamine;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;10/Modified site: amidated carboxyl end (Asn) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-reterer C;Superfamily:
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PA0110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ś
                                                                                                                                                                                                                                                                                    hypertrehalosemic/adipokinetic hormone - bollworm
N;Alternate names: Hez-HrTH
C;Species: Helicthis zea (bollworm, corn earworm, tomato fruitworm)
C;Date: 30-Jun-1989 #sequence_revision 23-Mar-1995 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-14 <SAV>
                                                                                                                                                                                                                                             C;Accession: A31571
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Bird,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
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A; Residues: 1-15 < KAM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Accession: PA0110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cross-references: UNIPROT:Q9SCX3; UNIPARC:UPI000017B01C Superfamily: translation elongation factor eEF-1 beta cl
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1995
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Pred. No. 1.4e
0; Mismatches
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N;Alternate names: Hez-AKH
C;Species: Heliothis zea (bollworm, corn earworm, t
C;Date: 31-Mar-1988 #sequence_revision 23-Mar-1995
C;Accession: A2424
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Gaede, G.; Wilps, H.; Kellner, R.
Biochem. J. 269, 309-313, 1990
A;Title: Isolation and structure of a novel charged member
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               adipokinetic hormone - nestling-sucking blowfly (C;Species: Protophormia terraenovae (nestling-sucking blowfly) C;Date: 17-Apr:1993 #sequence_revision 17-Apr:1993 #text_change 09-Jul-2004 C;Accession: S:1545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: protein
A;Residues: 1-11 <BUR>
A;Residues: UNIPROT:Q7MIH1; UNIPARC:UPI000017B0BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEBS Lett 339, 139-141, 1994
A;Title: Identification of a chaperonin-10 homologue in A;Reference number: S41747; MUID:94148071; PMID:7906228
A;Accession: S41747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Solanum tuberosum (potato)
C;Date: 19-Mar-1997 #sequence_revision 29-Aug-1997 #text_change 09-Jul-2004
C;Accession: S41747
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A; Residues: 1-8 < GAE>
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                                                                                          adipokinetic hormone - bollworm
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Pred. No. 1.4e+03
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                                                          tomato fruitworm)
                                 #text_change 09-Jul-2004
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B.A.; Holman, G.M.; Wagner,

R.M.;

Ridgwa

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Biochem. Biophys. Res. Commun. 135, 622-628, 1986
A;Title: Isolation and primary structure of a peptide from the corpora cardi
A;Reference number: A24244; MUID:86186794; PMID:3964263
A;Accession: A24244
A;Molecule type: protein
A;Residues: 1-9 <JAF>
A;Rosidues: 1-9 <JAF>
A;Cross-references: UNIPROT:P08901; UNIPARC:UPI000017661D
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide;
F;1/Modified site: pyrrolidone carboxylc acid (Gln) #status experimental
F;9/Modified site: amidated carboxyl end (Gly) #status experimental
                                                                                                                                                                              R;Rao, M.; Khadilkar, S.; Bandivadekar, K.R.; Deshpande, Biochem. J. 316, 771-775, 1996
A;Title: Structural environment of an essential cysteine A;Reference number: S70615; MUID:96265041; PMID:8670151
A;Cross-references: UNIPARC:UPI000017AE36
A;Experimental source: Chainia sp. strain NCL
A;Note: the source is designated as Chainia sp
C;Function:
C;Function: endohydrolyzation of beta-1,4-x
A;Pathway: fermentation of hemicellulose into
                                                                                                                                                                                                                                                                  C;Species: Streptomyces sp.
A;Variety: Chainia sp. NCL 82.5.1
A;Date: 19-Mar-1998 #sequence_revision
C;Accession: S70615
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A; Residues: 1-5 < RAO>
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A;Molecule type: protein
A;Residues: 1-15 <GAL>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEBS Lett. 298, 33-35, 1992
A;Title: Phosphorylation of
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Matches 4
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Date: 19-Mar-1997 #sequence_revision 11-Jun-1999
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Pred. No.
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Pred. No. 3e+03;
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lymnaDFamide 1 - great pond snail
C;Species: Lymnaea stagnalis (great pond snail)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_ct
C;Accession: S32471
R;Johnsen, A.H.; Rehfeld, J.F.
Bur. J. Biochem. 213, 875-879, 1993
A;Title: LymnaDFamides, a new family of neuropeptides from A;Reference number: S32471; MUID:93238777; PMID:8477756
A;Accession: S32471
                                                          -, m...auramide 2 - great pond snail
C. Species: Lymnaea stagnalis (great poi
C. Date: 19-Mar-1997 #sequence_revision
C. Accession: $32472
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S32472
R;Jöhnsen, A.H.; Rehfeld, J.F.
Eur. J. Biochem. 213, 875-879, 1993
A;Title: LymnaDFamides, a new family of neuropeptides from A;Reference number: S32471; MUID:93238777; PMID:8477756
                                                                                                                                                                                                                                                                                                                                 A;Experimental source: ganglia C;Keywords: amidated carboxyl end; neuropeptide F;13/Modified site: amidated carboxyl end (Phe)
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A;Title: A novel class of plant proteins containing a h
A;Reference number: S16323; MUID:91266907; PMID:1675603
A;Accession: S16324
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 21-Nov-1993 #sequence_revision 12-May-1995 #text_change 21-Jul-2000
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                                                                                                                                                                                                                                                                                  Best
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;Residues: 1-8 <RUB>
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on 19-Mar-1997
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lymnaDFamide 5 - great pond snail
C;Species: Lymnaea stagnalis (great pond snail)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
C;Accession: S32475
R;Johnsen, A.H.; Rehfeld, J.F.
Bur. J. Biochem. 213, 875-879, 1993
Bur. J. Biochem. 213, 875-879, 1993
A;Title: LymnaDFamides, a new family of neuropeptides from the pond snail, 1
A;Reference number: S32471; MUID:93238777; PMID:8477756
A;Accession: S32475
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A;Molecule type: protein
A;Residues: 1-13 <JOH>
A;Cross-references: UNIPROT:P80179; UNIPARC:UPI00001303CA; PIDN:AAB26363.1; PID:g299830
A;Experimental source: ganglia
C;Keywords: amidated carboxyl end; neuropeptide
F;13/Modified site: amidated carboxyl end (Phe) #status predicted
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Eur. J. Biochem. 213, 875-879, 1193
Eur. J. Biochem. 213, 875-879, 1193
A;Title: LymnaDFamides, a new family of neuropeptides from the pond snail, Lymnaea stagn A;Reference number: S32471; MUID:93238777; PMID:8477756
A;Accession: S32474
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S32474
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A;Residues: 1-13 <JOH>
A;Cross-references: UNIPROT:P80182; UNIPARC:UPI00001303D2; PIDN:AAB26366.1; PID:g299833
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Search completed: July Job time : 39 secs
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Matches 4
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;Species: Lymnaea stagnalis (great pond snail)
;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
;Accession: S32474
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Pred. No. 3.6e+03;
1; Mismatches 3
                       23:00:04
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Pred. No. 3.6e+03;
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       Copyright
       GenCore version 5.1.9 (c) 1993 - 2006 Biocceleration Ltd.
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RESULT 2
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8 CRC64; ; Length 1 ; 4; Indel	AA. DNA stage; Flagroup. DNA stage; Flagroup. Group. Group	
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RESULT 3
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ID Q9BGG8;
DT 01-JUN-2
DT 01-JUN-2
DT 07-FEB-2
DT Thyroid
GN Name=THR
OS Sorex ar
OC Eukaryot
OC Mammalia
OC Mommalia
OC NCBI_Tax
RN [1]
RA Larkin I
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R InterPro; IPR000886; ER_target_S.

R PROSITE; PS00803; CALRETICULIN_1; PARTIAL.

DR PROSITE; PS00805; CALRETICULIN_REPEAT; PARTIAL.

DR PROSITE; PS00805; CALRETICULIN_REPEAT; PARTIAL.

DR PROSITE; PS00804; ER_TARGET; PARTIAL.

RW Calcium; Chaperone; Direct protein sequencing; Endoplasmic retirection; Metal-binding; Zinc.

Calreticulin.

Calreticulin.

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Calreticulin.
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Am. J. Trop. Med. Hyg. 67:636-639(2002).

-!- FUNCTION: Molecular calcium binding chaperone promotinoligomeric assembly and quality control in the ER via calreticulin/calnexin cycle. This lectin may interact
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01-OCT-2001, sequence versi
07-FEB-2006, entry version
Calreticulin (Fragment).
                                                                                                                                                                                                                                                                          01-JUN-2001, integrated into Unil
01-JUN-2001, sequence version 1.
07-FEB-2006, entry version 9.
Thyroid hormone receptor alpha ()
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Eukaryota; Entamoebidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/taDistributed under the Creative Commons Attribution-NoDerivs License
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STRAIN=HM-1:IMSS;
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                            NUCLEOTIDE
                                                                                          NCBI_TaxID=42254;
                                                                                                                                                         Mammalia;
                                                                                                                                                                                                                  Sorex araneus (Eurasian
                                                                                                                                                                                                                                                    Name=THRA1;
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GO:0005788; C:endoplasmic reticulum lumen;
GO:000509; F:cellcium ion binding; ISS.
GO:000509; F:cellcium ion homeostasis; ISS.
GO:0006874; F:calcium ion homeostasis; ISS.
GO:0006811; P:protein-nucleus export; ISS.
GO:0006355; P:regulation of transcription,
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SUBCELLULAR LOCATION: Endoplasmic reticulum lumen (By similari
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         Zhdanova N.;
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No. 1.8e+03;
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RESULT 4
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ID Q6SE52_D
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Q9XNQ1 BOOMI
ID Q9XNQ1 BOOMI PRELIMINARY; PRT; 17 AA.

AC Q9XNQ1;
DT 01-NOV-1999, integrated into UniProtKB/TrEMBL
DT 01-NOV-1999, sequence version 1.

DT 01-FEB-2006, entry version 10.
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Q6SE52;
05-JUL-2004,
                                                   Cytochrome oxidase subunit 2 (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PubMed=14762063; DOI=10.1101/gr.1329204;
Halligan D.L., Eyre-Walker A., Andolfatto P., Keis
"Patterns of evolutionary constraints in intronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Drosophila simulans (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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                Boophilus microplus Mitochondrion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of Drosophila.";
Genome Res. 14:273-279(2004).
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07-FEB-2006, entry version
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Eukaryota; Metazoa;
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Arthropoda; Chelicerata; Arachnida; Acari;
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83.3%;
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ivity; IEA.
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EMBL;
EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=95014476; PubMed=7929358; Ahuja S.K., Shetty A., Tiffany H.L., Murphy "Comparison of the genomic organization and human interleukin-8 receptors A and B.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JUL-2004, integrated into U
05-JUL-2004, sequence version
07-FEB-2006, entry version 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q6LCZ7_HUMAN
Q6LCZ7;
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                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens (Human)
Eukaryota; Metazoa;
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MEDLINE=99297341; PubMed=10368952;
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NCBI_TaxID=6941;
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Mammalia; Eutheria;
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U11874; AAA64382.1; -; mRNA.
U11875; AAA64383.1; -; mRNA.
U11876; AAA64384.1; -; mRNA.
U11877; AAA64386.1; -; mRNA.
U11878; AAA64386.1; -; mRNA.
U11872; AAA64380.1; -; mRNA.
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                                                            LSQETFSDLW 12
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                                                                                                                         Score 28; DB 2; Pred. No. 5.8e+03; Mismatches 4
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Pred. No. 4
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Q2QKW4;
Q2QKW4;
24-JAN-2006, integrated into UniProtKB/TrEMBL.
24-JAN-2006, sequence version 1.
21-FEB-2006, entry version 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Paraneoptera; Hemiptera; Sternorrhyncha;
Aphidoidea; Aphididae; Macrosiphini; Hyadaphis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JAN-2006, sequence version 3. 21-FEB-2006, entry version 3. Cytochrome c oxidase subunit
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Q5SR29;
21-DEC-2004, integrated into Unil
21-DEC-2004, sequence version 1.
07-FEB-2006, entry version 7.
Hepatic leukemia factor (Fragment
                                                                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprobistributed under the Creative Commons Attribution-NoDerivs
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PubMed=16368250;
von Dohlen C.D., Rowe C.A., Heie O.E.;
"A test of morphological hypotheses for tribal and subtribal
relationships of Aphidinae (Insecta: Hemiptera: Aphididae) u
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Mammalia; Eutheria; Euarchontoglires; G
Muroidea; Muridae; Murinae; Mus.
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Name=Hlf; ORFNames=RP23-220F20.3-004;
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von Dohlen C.D., Rowe C.A., Heie O.E.;
von Dohlen C.D. Rowe C.A., Heie O.E.;
"A test of morphological hypotheses for tribal and subtriba
relationships of Aphidinae (Insecta: Hemiptera: Aphididae)
sequences.";
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24-JAN-2006, sequence version 1.
21-FEB-2006, entry version 3.
Cytochrome c oxidase subunit I (
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Neoptera; Paraneoptera; Hemiptera; Sternorrhyncha;
Aphidoidea; Aphididae; Hyalopterus.
NCBI_TaxID=312888;
                                                                                                  Wada N., Matsumura
Nakanishi Y.;
                                                                                                                         MEDLINE=95355401; PubMed=7543095; Wada N., Matsumura M., Ohba Y., Ko
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Mammalia; Eutheria;
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Commons Attribution-NoDerivs License
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Kobayashi
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NUCLEOTIDE SEQUENCE.
Muschen M., Re D., Br.
Kuppers R., Rajewsky i
Submitted (SEP-2000)
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EMBL; A.
EMBL; A.
NON TER
                                                                                                                                                                                                                                                                                                                         PubMed=16343953; DOI=10.1016/j.ympev.2005.10.003; Larkin L.L., Neff J.L., Simpson B.B.; Larkin L.L., Neff J.L., Simpson B.B.; "Phylogeny of the Callandrena subgenus of Andrena Andrenidae) based on mitochondrial and nuclear DN convergent evolution.";
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Q71H30;
Q71H30;
Q71H30;
Q7-JUL-2004, integrated into UniProtKB/
Q5-JUL-2004, sequence version 1.
Q7-MAR-2006, entry version 9.
Cytochrome oxidase subunit I (Fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neoptera; Endopterygota;
Andrenidae; Andreninae;
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Eur. J. Immunol. 32:3785-3792(2002).
          SEQUENCE
                                     Mitochondrion.
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Distributed under the Creative
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AJ279012; CAC35540.1; -; Genomic_DNA.
AJ279013; CAC35541.1; -; Genomic_DNA.
AJ279013; CAC35541.1; -; Genomic_DNA.
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Endopterygota; Hymenoptera; Apocrita; Aculeata; Apoidea;
; Andreninae; Andrena; Callandrena.
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PROTEIN SEQUENCE, SUBCELLULAR LOCATION, AN STRAIN=cv. DE GRACE; TISSUE=LBAF; MEDLINE=20181728; PubMed=10715320; DOI=10. Peltler J.-B., Friso G., Kalume D.E., Roeg Adamska I., van Wijk K.J.; "Proteomics of the chloroplast: systematic targeting analysis of lumenal and peripher Plant Cell 12:319-341(2000).
                                                                                                                                                                     P82326;
01-JUN-2000, integrated into UniProtKB/TrEMBL.
01-JUN-2000, sequence version 1.
07-FEB-2006, entry version 14.
Unknown protein from 2D-page of thylakoid lumen (SPOT107) (Fragment)
                                                                                                                              Pisum sativum (Garden pea).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicotyledons;
rosids; eurosids I; Fabales; Fabaceae; Papilionoideae; Vicieae; Pi
                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; UC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thuerauf D.J., Hanford D.S., Glembotski C.C., "Regulation of rat brain natriuretic peptide transcription. potential role for GATA-related transcription factors in my
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN-Sprague Dawley;
MEDLINE-94299479; PubMed-8027030;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi; Muroldea; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-2004, integrated int 05-JUL-2004, sequence version 07-FEB-2006, entry version
                                                                                                                                                                                                                                                         P82326_PEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                          NCBI_TaxID=3888;
                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Copyrighted by the UniProt Consortium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rattus norvegicus (Rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=BNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brain natriuretic peptide (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biol. Chem. 269:17772-17775(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene expression."
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ER 19 19
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Pred. No. 1.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genomic_DNA.
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Pred. No. 7.6e+03;
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                                     systematic
                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                         peripheral
                                                           DOI=10.1105/tpc.12.3.319;
E., Roepstorff P., Nilsson
 THYLAKOID MEMBRANE LUMEN
                                                                                                                                                                                                                                                       14
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al thylakoid protei
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RESULT 15
Q9UJZ3 HUMAN
ID Q9UJZ3;
AC Q9UJZ3;
DT 01-MAY-2
DT 07-FEB-2
DT 07-FEB-2
DE Oviduct
GN Name=CGB
OS Homo sap
OC Eukaryot
OC Mammalia
OC Homo.
OX NCBI_Tax
RN [1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 14
Q9QVS7
ID Q9QVS
AC Q9QVS
AC Q9QVS
AC Q9QVS
DT 01-WA
DT 07-FE
DE Lacta
OS Mus s
OC Eukaa
OC Muroi
OX NCBI.
RN (1)
RN (1)
RP NUCLE
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Best Local :
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Best Local (
                                                                                                                                                                                                                 Q9UJZ3 HUMAN
Q9UJZ3;
01-MAY-2000,
  Homo.
NCBI_TaxID=9606;
[1]
                                                                                                                                                                      01-MAY-2000,
07-FEB-2006,
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Q9QVS7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GO; GO:0009507; C:chloro
GO; GO:0009579; C:thylal
Chloroplast; Thylakoid.
NON_TER 14 14
                                                                                       Eukaryota;
                                                                                                            Homo sapiens (Human)
                                                                                                                                Name=OGP
                                                                                                                                                  Oviduct glycoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
MEDLINE=95201434; PubMed=7534515;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria; Euarchor
Muroidea; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus sp.
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07-FEB-2006, entry version 11.
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                                                                 Mammalia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10095;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <del>-</del>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DEVELOPMENTAL STAGE: UNFOLDED AND FULLY DEVELOPED LEAVES MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UN PROTEIN IS: 6.0, ITS MW IS: 18.3 kDa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GO:0009507; C:chloroplast; IEA.
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                                                            .; Metazoa;
Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                      sequence version entry version 10
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                                                               Chordata; Craniata; Vertebr
Euarchontoglires; Primates;
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30.8%;
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83.3%;
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                                                            Vertebrata; Euteleo
rimates; Catarrhini;
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                                                                                   Euteleostomi;
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GenCore version 5.1.9
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Minimum
Maximum
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Maximum Match
Listing first
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Perfect score:
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A_Geneseq_8:*
1: geneseqp198
2: geneseqp200
4: geneseqp200
5: geneseqp200
6: geneseqp200
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9: geneseqp200
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Gapop 10.0 ,
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1 PPLSQETFSDLWKLLPENG 19
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             geneseqp2004s:*
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45 gt
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Pred. No: is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

2222233	Result No.
105 97 97 92 92 92 92 92 92 92 93 81 81 81 77 77 77	Score 105
100.0 92.4 92.4 87.6 87.6 81.0 81.0 81.0 81.0 77.1 77.1 77.1 77.1 77.1 77.1 77.1 7	Query Query Match Length
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AAW37238 AAW47074 AAW47074 AAB230870 ABG73435 ADR20234 AAB76310 ABF16088 AAY66310 ABF75806 AAY66310 ABG75806 ADC2252 ADG7883 ADD49951 ABD9951 ABD9951 ABB21960 AAR54910 ABB7960 AAR54910 AAR54910 AAR54911 AAB29157 AAB89914 AAB899130 AAR6899130 AAR6899130	3 ID
Aaw92319 p53 N-ter Aaw47074 p53/RB in Aaw47074 p53/RB in Aaw47074 p53/RB in Aae30870 Peptide u Abg73435 Human p53 Adr20234 Mixed ele Aae30872 BOX-1 dom Ae616088 Novel sta Aay06310 Human p53 Abg75806 Solid pha Adc22292 Protein pha Adc22292 Protein pha Adg78883 Human p53 Adg4883 Human p53 Adg480537 Peptide # Adg00921 Pancreatic Aed80637 Human p53 Aee74960 p53/MIM2- Aar54909 Immunodom Aar54910 Immunodom Aar54910 Immunodom Aar54910 Imgurodom Aar54910 p53 prote Aab29157 Peptide # Aag89730 p53 DR 3a Aag89500 p53 DR 3a	Description Aaw82321 p53 homol

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65.7	66.7	66.7	66.7	67.6	67.6	67.6	69.5	69.5	69.5	69.5	69.5	69.5	70.5	70.5	70.5	70.5	71.4	73.3	73.3	73.3	73.3
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AAB29162	ADT78987	ADT79007	AAY57799	AAB29160	AAB29161	AAY99001	AEF16090	ADT78973	AAE30860	AAB29167	ABB05528	ADZ38449	ADT79028	AAB29163	AAB29159	ADU00102	AEF16089	712	522	ADT02875	ABG73434
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162					161	001	6090	973	860	167	528	449	028	163	159	102	5809	122	227	875	434
Peptide	DNA	Human p5	TRAM-int	Peptide	Peptide	HLA class	No	Human p53	EGF	Pep	Biotinyla	Human kir	Mou	Pep	Peptide #	Ami	No	Human p53	p 53	Human	Human
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ALIGNMENTS

AAW82321 ID AAW8

AAW82321 standard; peptide; 19

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RESULT 1

Synthetic. 22-APR-1997; 20-APR-1998; 29-OCT-1998. WO9847919-A1 p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament. p53 homologue TIP peptide. 22-FEB-1999 AAW82321; (UYDU-) UNIV DUNDEE. (first entry) 97GB-00008089 98WO-GB001140

Lane DP;

WPI; 1998-609975/51.

New substance with a mdm2 binding domain and coupling partner - useful for stabilising in cells without an efficient mdm2-mediated degradation pathway. degradation

Disclosure; Fig 1; 52pp; English.

This sequence is a peptide homologue of a region of p53 which binds to mdm2. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows level p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 function. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53

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RESULT 2
AAW82319
ID AAW8
XX AAW
AC AAW8
XX P53
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                                                                               RESULT 3
AAW37228
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Best Local Similarity
Matches 19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence is a peptide homologue of a region of p53 which binds to mdm2. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 function. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with
  AAW37228;
                                                       AAW37228 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New agents which inhibit interaction of p53 and mdm2 - useful for activating p53, e.g. for treating cancers, viral conditions or other conditions associated with non functional p53 or mdm2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 non-functional p53
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                                                    peptide; 18
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                                                                                                                                                                                                                                                                                                                                  Score 105; DB 2;
Pred. No. 1.7e-09;
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Pred. No. 1.7e-09;
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p53/RB interaction inhibiting peptide 6 (residues 11-29

of human p53).

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RESULT 4
AAW47074
ID AAW4
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                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Compounds binding to MDM2 protein and inhibit its interaction with p53 useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 34; 45pp; English.
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07-APR-1997;
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     19-MAY-1998
                                                                                                       AAW47074 standard;
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                                                                                                                                                                                                                                                                                                                                             Similarity 100.
18; Conservative
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(first entry)
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97GB-00007041.
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                                                                                                       peptide;
                                                                                                                                                                                                                                                                                                                                  94.3%; Jr
100.0%; Pr
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Chene P, Furet P;
                                                                                                          19
                                                                                                                                                                                                                                                                                                                                             Score 99; DB; Pred. No. 1.5
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hes 0;
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Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This peptide fragment of p53 is an inhibitor of the interaction between a p53 protein and a retinoblascoma (RB) protein. This peptide corresponds to residues 11-29 of human p53 (AMA47079). The interaction between p53 and Rb is found to be critical for determining whether or not a cell enters apoptosis. Apoptosis is prevented if interaction occurs. The interaction is between regions 1-71 or 290-393 of p53 and region 379-928 of Rb. The invention provides methods to identify compounds able to modulate interaction or binding between p53 and Rb protein. The method comprises combining p53 and Rb, or their fragments, with a test compound and detecting interaction/binding between p53 and Rb protein. The method and detecting interaction/binding between them. These inhibitory compounds are used in human or veterinary medicine to modulate p53 activity and processes, specifically for inducing apoptosis in tumour cells (possibly also in cells infected by virus), in vivo or in vitro. Expression of these modulators by gene therapy methods is also contemplated. Other activities that can be affected are transcription repression, G1 arrest, DNA repair, homologous recombination and 3'-5'-exonuclease activity. Modulation of interaction with Rb may also
                                                                                                                                                           24-FEB-2003
                                                                                                                                                                                                                       AAE30870 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 AA;
                                                                           p53 polypeptide; p
cancer; ischaemia;
                                                                                                                        Peptide used in modulating the binding of p53 to p300.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying compounds that modulate interaction of p53 and Rb protein or those that bind to Rb protein, used to induce apoptosis, specifical for treatment of tumours.
               WO200265134-A2
                                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Retinoblastoma gene; RB; p53 protein; interaction; inhibitor; tumour; apoptosis; modulator; medicine; veterinary; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CANC-) CANCER
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17; Conserv
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                                                                                                                                                                                                                                                                                                      PPLSQETFSDLWKLIPEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 63; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                         (first entry)
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                                                                           p300 polypeptide; cell cycle; cell death; a; cytostatic; vasotropic.
                                                                                                                                                                                                                       peptide; 19
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Pred. No. 3.3e-08;
1; Mismatches 0
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RESULT 6
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Best Local &
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Novel polypeptide that inhibits degradation of endogenous mammalian cell, useful for treating cancer, e.g. cervical tumor such as sarcoma or carcinoma.
                                                                                                                                                                                                                                                                             carcinoma; squamous cell carcinoma; lung cancer; pancreatic cancer
leukaemia; lymphoma; neuroblastoma; sarcoma; osteosarcoma; gliobla;
colon carcinoma; melanoma; choriocarcinoma; breast carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a peptide for use in modulating the binding of a p53 polypeptide to a p300 polypeptide. The new peptide is useful in modulating the binding of a p53 polypeptide to a p300 polypeptide. The peptide may be used to regulate the mammalian cell cycle or to induce or prevent cell death, for the treatment of cancer or ischaemia. The invention is useful in gene therapy. The present sequence is a peptide used in modulating the binding of p53 to p300
                                                                                                                                                                                                                                                                                                       Human; p53; degradation; cervical cancer; tumour; ovarian cancer; glioma; carcinoma; squamous cell carcinoma; lung cancer; pancreatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New peptide for modulating the binding of p53 polypeptide to p300 polypeptide, useful for regulating the mammalian cell cycle for the treatment of cancer or ischemia.
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                                                                         Yuan Z,
                                                                                                (YUAN/) YUAN Z. (GUJJ/) GU J.
                                                                                                                                       08-DEC-1999;
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                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                           Human p53 peptide fragment
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                                                                                                                                                                                                                                                                                                                                                                                                                        ABG73435 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                       peptide; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90.5%;
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           p53 in cancer
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Matches 17
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The invention describes a composition (I) for assessing the presence of at least a first target molecule in a sample, comprising: several low-to moderate affinity binding elements distributed on a surface of, and operatively coupled to a support, where concomitant binding of the first target molecule to two or more of the binding elements results in a high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a substantially pure polypeptide that inhibits degradation of endogenous p53 in a mammalian cell. The polypeptide or a synthetic polypeptide comprising a region of the pure polypeptide is useful for inhibiting degradation of endogenous p53 in a mammalian cell.
                                                                                                                                                                                                                                                         as polypeptide in sample e.g., blood, comprises several low-to-modern affinity binding elements distributed on surface of, and operatively coupled to support.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
                                                                                                                                                                                                                                                                                                                           Composition useful for assessing presence of first target molecule such as polypeptide in sample e.g., blood, comprises several low-to-moderate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnostic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mixed element capture agent study associated peptide Gal80bp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-NOV-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9.9
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                                                                                                                                                                                                Example 2; SEQ ID NO 9;
                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-593073/57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KODA/) KODADEK T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-JAN-2003; 2003US-0438805P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     target molecule
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      biological pathway; protein detection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     detection; binding element; concomitant binding;
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100.0%; Pr
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                                                                                                                                                                                                58pp; English
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hes 0;
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                                              New peptide for modulating the binding of p53 polypeptide to p300 polypeptide, useful for regulating the mammalian cell cycle for titreatment of cancer or ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 p53 polypeptide; p300 polypeptide; cell cycle; cell death; gene therapy; cancer; ischaemia; cytostatic; vasotropic.
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100.0%; Pred. No. 2.
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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Example 2; Fig 6C; 87pp; English

¥868888888

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RESULT 9
AEF16088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAY-1999;
26-NOV-1999;
18-MAY-2000;
This invention relates to a novel stabilized alpha-helix peptide structure which comprises a selected number of any combination of natural and unnatural amino acids (where the peptide further comprises at least one cross-linking moiety that stabilizes the peptide in alpha-helix conformation). The invention may be useful for the development of compounds with a cytostatic or virucide activity acting as a cell proliferation inhibitor, viral replication inhibitor, gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a peptide for use in modulating the binding of a p53 polypeptide to a p300 polypeptide. The new peptide is useful in modulating the binding of a p53 polypeptide to a p300 polypeptide. The peptide may be used to regulate the mammalian cell cycle or to induce or prevent cell death, for the treatment of cancer or ischaemia. The invention is useful in gene therapy. The present sequence is BOX-1 domain
                                                                                                                                                                                                              New stabilized alpha-helix peptide structure useful in medicinal chemistry as therapeutic agents comprises selected number of any combination of natural and unnatural amino acids, where the peptifurther comprises cross-linking moieties.
                                                                                                                                                                          Disclosure;
                                                                                                                                                                                                                                                                                                                          WPI; 2006-088591/09
                                                                                                                                                                                                                                                                                                                                                                                                            (VERD/)
(SCHA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-JUN-2005; 2005US-00148976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-JAN-2006
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein structure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel stabilized
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  replication;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
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                                                                                                                                                                                                                                                                                                                                                                                                          VERDINE G L.
SCHAFMEISTER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLSQETFSDLWKLLP 16
                                                                                                                                                                          Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A
A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   p300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    83.8%; Silarity 100.0%; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-00574086
                                                                                                                                                                                                                                                                                                                                                                    Schafmeister
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0134708P.
99US-0167634P.
                                                                                                                                                                      16; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            alpha-helix
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic; virucide; gene expression; cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide; 16
                                                                                                                                                                                                                                                                                                                                                                                                            a
                                                                                                                                                                                                                                                                                                                                                                                                     [F]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "N-terminal acetyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 88;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B
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                                                                                                                                                                                                                                    amino acids, where the peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              structure-related peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6; Le
7.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cell proliferation;
signaling.
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Query Match

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Sequence 15

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RESULT 10
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ពួកក្រកួតក្រកួតក្រកួត
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        other cell signaling process inhibitor. The stabilized alpha-helix peptide structure generates stabilized-helical structures, as well as other secondary structures, to enable the investigation of complex structure-function relationships in proteins and enable the development of new therapeutics incorporating specific stabilized secondary structure motifs. The peptides can be synthesized one-at-a time, using a traditional peptide to generate the structure motif, and may control (promote or inhibit) cell functions. The present sequence is that of a peptide which was used during the exemplification of the novel stabilized alpha-helix peptide structures of the invention.
                                       This synthetic peptide corresponds to amino acids 13-27 of human p53. It was used to raise anti-peptide p53.1 and anti-RB18A antibodies in rabbit. RB18A (see AAY06309) is a novel p53 regulatory protein of the invention. The antibodies were used in immunoscreenings that led to the isolation of a CDNA clone (see AAX59124) encoding human RB18A. The invention provides methods and compositions for the diagnostic and therapeutic applications of RB18A, in particular for the diagnosis, prevention or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             wound healing; graft rejection; reperfusion injury; wound healing; graft rejection; reperfusion injury; myocardial infarction; stroke; traumatic brain injury; neurodegenerative disease; ischaemia; toxaemia; infection; neurodegenerative disease; ovarian cancer; colon cancer; di
                                                                                                                                                                                                         New p53 regulatory protein (RB18A) useful primers to detect the transcription rate a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16 AA;
                          neoplasia
                                                                                                                                                               Example 1;
                                                                                                                                                                                                                                                        WPI; 1999-395177/33.
                                                                                                                                                                                                                                                                                     Frade
                                                                                                                                                                                                                                                                                                                                               15-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                             14-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                           24-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9931231-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RB18A; p53 regulatory protein; apoptosis; neoplasia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY06310;
                                                                                                                                                                                                                                                                                                                 (INRM ) INSERM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY06310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        p53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LSQETFSDLWKLLPEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LSQETFSDLWKLLPEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide (aa13-27).
                                                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                                                                                 INST NAT SANTE & RECH MEDICALE
                                                                                                                                                                                                                                                                                                                                               97EP-00403051.
                                                                                                                                                                                                                                                                                                                                                                             98WO-EP008560
                                                                                                                                                                25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide;
                                                                                                                                                            87pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81.0%;
100.0%;
                                                                                                                                                               English
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Pred. No.
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2.3e-06;
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and
                                                                                                                                                                                                            abundance
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nnce of RB18A mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inflammation;
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in
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RESULT 11
ABG75806
Best Local Similarity
Matches 15; Conserv
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Isolation of biological and/or organic solvent,
                                                                                                                                                                       Modified-site
                                                                                                                                                                                                 Modified-site
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                                                                                                                                                                                                                                                                              Modified-site
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                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                   Solid phase
                                                                                                                                                                                                                                                                                                                                                                                                                 08-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                            ABG75806 standard; peptide;
                     WPI; 2003-298114/29.
                                                 (DULY/)
(LAUR/)
                                                             (UDEL/)
                                                                                          07-JUL-1999;
                                                                                                       06-JUL-2000; 2000US-00610761
                                                                                                                      12-NOV-2002
                                                                                                                                    US6479296-B1
                                                                                                                                                                                                                                                                                                        Modified-site
                                                                            (BONN/)
                                                      ) UDELL L S.
) CREASEY W A.
) DULY S.
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                                  Æ,
                                                 LAURSEN R A.
                                                                            BONNER A G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                  Udell LS,
                                                                                                                                                                                                                                                                                                                                                                                                  precipitation and extraction method peptidyl-resin
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                                                                                          99US-0142582P
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specification)"
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specification)
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specification)"
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'note= "OTHER=
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                                                                                                                                                                                                                                                                                                                        "OTHER=
                                                                                                                                                                                                         "OTHER=
                                  Creasey
analyte, e.g. protein, from sample, e.g. aqueous involves precipitating biological analyte on
                                                                                                                                                              OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 9.3e-06; ; Mismatches 0;
                                  WA,
                                                                                                                                                  modified with Trt and linked to the
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Tentagel resin"
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                                                                                                                                                                                                                         CC analyte (e.g. a protein and/or nucleic acid from a sample or aqueous and/or organic solvent) which comprises the precipitation of the composition of the precipitated biological analyte off the composition and extraction (SPE) method. The method can be used in composition and extraction (SPE) method. The method can be used in composition and extraction (SPE) method. The method can be used in composition and in the purification of combinatorially-composition and in the purification of combinatorially-composition and composition and in the purification of combinatorially-composition and composition and manipulation of milligram amounts of precipitate. The method can be automated without difficulty and provides convenience for multiple composition and manipulation of milligram amounts of precipitate. The method composition and method without difficulty and provides convenience for multiple composition and manipulation of method composition and manipulation of milligram amounts of precipitate. The method composition and manipulation of milligram amounts of precipitate. The method composition and manipulation of milligram amounts of precipitate. The method composition and manipulation of milligram amounts of precipitate. The method composition and manipulation of milligram amounts of precipitate. The method composition and method composition and method without difficulty and provides convenience for multiple composition and method composition and method composition and method composition and compo
Matches
                                Query Match
Best Local
                                                                                                                                      Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            solid-phase extraction device, and eluting precipitated analyte from
15;
                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   device
                                                                                                                                         A,
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            discloses
                                77.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a method for the isolation of a biological
0,
                                Score 81;
Pred. No.
   Mismatches
   DB 6; Le
9.3e-06;
hes 0;
                                                                 Length 15;
       Indels
       0;
       Gaps
       0;
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RESULT 12
ADC22292
                                                                                                       01-AUG-2001; 2001US-0309395P
13-DEC-2001; 2001US-0341589P
                                                                                                                                                                                                                                                                              localisation domain; binding domain; subcellular compartment localisation
                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC22292 standard;
 N-PSDB;
                                             Bright G,
                                                                                                                                                      01-AUG-2002; 2002WO-US024572
                                                                                                                                                                                      13-FEB-2003
                                                                                                                                                                                                                    WO2003012068-A2
                                                                                                                                                                                                                                                                                                                                          Protein binding domain amino acid sequence SEQ ID NO:141.
                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                             recombinant fusion protein; fusion protein;
                                                                                                                                                                                                                                                                                                                                                                         18-DEC-2003 (first entry)
2003-248174/24.
DB; ADC22293.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 SQETFSDLWKLLPEN
                                                                            CELLOMICS INC.
                                             Premkumar
                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide;
                                             DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15
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                                              Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                        15
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                                                                                                                                                                                                                                                                                                              binding; detection;
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New recombinant fusion protein comprising detection and first localization domains and a binding domain for the molecule of useful for detecting binding of a molecule of interest.

interest,

Claim 8; SEQ ID NO 141; 101pp; English

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RESULT 13
ADG78883
ID ADG78
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Best Local (
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The present sequence is that of a peptide (PNC27) comprising amino aci residues 12-26 of human p53. The invention provides peptides corresponding to all or a portion ADG78884-ADG78885 of this sequence, peptides being lethal to malignant or transformed cells when fused to membrane-penetrating leader sequence. The peptides are thus useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; p53; cytostatic.
                                                                                                                                                                                                          Claim 2; SEQ ID NO 1; 32pp; English.
                                                                                                                                                                                                                                                                                                                    New peptides useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-MAR-2002; 2002US-0363785P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human p53 protein peptide PNC27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYNY ) UNIV NEW YORK STATE RES FOUND (PINC/) PINCUS M R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-MAR-2003; 2003WO-US007687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG78883 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present
                                                                                                                                                                                                                                                                            sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cells. The present sequence is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PPLSQETFSDLWKLL 15
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                                                                                                                                                                                                                                                                            of cell cycle
                                                                                                                                                                                                                                                                            for treatment of neoplastic diseases comprises ll cycle regulatory protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 81;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (residues 12-26).
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RESULT 14
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    penetrating leader sequence. The peptides are derived from the human p53 composition. Also disclosed are (i) a pharmaceutical composition comprising at least one of the peptides or its analogues or derivatives admixed with a pharmaceutical carrier, and (ii) a method of selectively killing malignant or neoplastic cells in a subject. The leader sequence is preferably located at the carboxy terminal end of the peptide, its analogue or derivative. The leader sequence comprises predominantly positively charged amino acid residues. The leader sequence is at least one of penetratin, Args, TAT of HIVI, D-TAT, R-TAT, SV40-NLS, CCMV GAG, p22N, Lambda N, Delta N, yeast PRP6, human U2AF, human C-FOS, human C-JUN, yeast GCN4 or p-vec. Selectively killing malignant or neoplastic cells in a subject comprises administering to the subject an amount of the peptide, where a membrane-penetrating leader sequence is fused to the carboxy terminal of the peptide, its analogue or derivative. The present
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Best Local (
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05-APR-2001; 2001US-00827683
12-MAR-2002; 2002US-0363785P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     treating neoplastic disease in an animal, preferably a human. Claimed methods of selectively killing malignant or neoplastic cells comprise administering the p53-derived peptide fused at its C-terminus to a membrane-penetrating leader sequence, such as the penetratin leader
                                                                                                                                                                                                                                            The present invention relates to peptides that are selectively lethal malignant and transformed mammalian cells when fused to a membrane-
                                                                                                                                                                                                                                                                                                                                        New peptide fused to membrane-penetrating leader sequence and selectively lethal to malignant or transformed cells, useful 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      membrane-penetrating sequence ADG78886.
                                                                                                                                                                                                                                                                                               Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                        Pincus MR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lethal peptide; malignant cell; transformed cell;
membrane-penetrating leader sequence; human; p53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15
                                                                                                                                                                                                                                                                                                                           neoplastic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2004038902-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neoplastic cell; cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADN48957 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PINC/) PINCUS M R.
                                                                                                                                                                                                                                                                                            2; SEQ ID NO 1;
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                                                                                                                                                                                                                                                                                                              or malignant cells, e.g. cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        from
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human p53 protein.
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                                                                                                                                                                                                                                                                                            9pp; English.
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J. 9.3e-06;
O;
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           isolatable from sera and other bodily fluids of pancreatic cancer captioners; (2) an immunologically active fragment of the pancreatic carcinoma-specific antigen 3C4-Ag; (3) an antibody or its binding portion having specificity to pancreatic carcinoma specific antigen 3C4-Ag; (4) a murine hybridoma cell line that produces a monoclonal antibody specifically immunoreactive with the 3C4-Ag antigen; (5) a monoclonal antibody, mb3C4, secreted by the hybridoma cell line; (6) detecting a cell, tissue, or fluid sample from the subject with (the binding portion of) an antibody that binds to 3C4-Ag; (b) detecting antibody-antigen complex in the sample; and (c) correlating the detection of elevated complex in the sample; and (c) correlating the detection of elevated complex of antibody-antigen complex in the sample with the presence of pancreatic cancer; (7) a diagnostic kit for detecting 3C4-Ag in a cell, tissue or fluid sample from a patient comprising; (a) the antibody or its binding portion specific for 3C4-Ag or its immunologically active fragment; (b) a conjugate of a specific binding partner for the antibody.
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                           3C4-Ag in substantially purified form, which has a molecular weight of 4.5 ADA as determined by SDS-PAGE, and a pI on isoelectrofocusing of 4.5 -5.0. The antigen is unglycosylated or minimally glycosylated and is primarily localised on the surface of rat and human pancreatic cancer cells but is not detected in normal, non-proliferating cells. Also described: (1) a soluble pancreatic carcinoma-specific antigen 3C4-Ag having a molecular weight of 36-38 kDa as determined by SDS-PAGE which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New pancreatic carcinoma-specific antigen 3C4-Ag primarily localized on
the surface of rat and human pancreatic cancer cells, but not detected
normal, non-proliferating cells, useful in diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2004065547-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Michl J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-OCT-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pancreatic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes the pancreatic carcinoma-specific antigen
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   binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bradu SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 1; 112pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           carcinoma-specific antigen 3C4-Ag; pancreatic cancer;
cytostatic; immunotoxin; pancreatic cancer associated antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer.
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   portion; and (c) a label
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               77.1%; Score 81; 100.0%; Pred. No.
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   for detecting the bound antibody
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                                                                                           Query Match
Best Local :
                                                                           Matches
                                                                                                                                                                                                    (8) treating pancreatic cancer in a patient by administering an effective amount of an antibody or its binding portion which specifically binds to 3C4-Ag or its immunologically active fragment, where the antibody or binding portion is conjugated or linked to a therapeutic drug or texin; and (9) a pharmaceutical composition comprising the antibody or its binding portion, and a carrier. The pancreatic carcinoma-specific antigen 3C4-Ag has cytostatic activity, and can be used as an immunotoxin. The antigen and methods are useful for detecting and treating pancreatic cancer. The present sequence represents a pancreatic cancer associated antigen peptide, which is used in the exemplification of the present
                                                                                                                                                  Sequence 15
                                                                                             Local Similarity
                                                                           15;
                                     1 PPLSQETFSDLWKLL
 PPLSQETFSDLWKLL 15
                                                                                                                                                  AA;
                                                                           Conservative
                                                                                             77.1%;
100.0%;
                                                                           0
                                                                                               Score 81;
Pred. No.
                                                                           Mismatches
                                                                                             DB 8; L
. 9.3e-06;
                                                                             0
                                                                                                              Length 15;
                                                                             Indels
                                                                             0,
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Search completed: July 5, 2006, 22:54:22 Job time : 194 secs

Run on:

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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
             seq length: 0
seq length: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query
Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           July 5, 2006, 23:00:16 ; Search time 181 Seconds (without alignments) 48.625 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BLOSUM62
Gapop 10.0 ,
             Published Applications AA Main:*

1 /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2 /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

3 /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US09_PUBCOMB.pep:*

4 /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

5 /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

5 /EMC_Celerra_SIDS3/ptodata/2/pubpaa/US11_PUBCOMB.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2097797 seqs, 463214858 residues
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105
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Copyright (c) 1993 - 2006 Biocceleration Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length DB
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                            US-09-214-371-74
US-09-72-262A-74
US-09-72-384-7
US-10-754-457-9
US-10-754-457-9
US-10-27-262A-1
US-10-27-262A-1
US-10-467-758-14
US-10-415-970-4
US-10-415-970-4
US-11-019-894A-1
US-11-019-894A-1
US-11-051-411-146
US-11-051-411-146
US-11-051-411-146
US-11-051-411-146
US-10-28-953-14
US-10-155-059-23
US-10-155-059-23
US-10-692-217-131
US-10-692-217-131
US-10-693-2388-131
US-10-632-388-131
US-10-651-723-143
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Sequence 74, Appl
Sequence 7, Appli
Sequence 7, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 14, Appli
Sequence 13, Appl
Sequence 1109, Ap
Sequence 1109, Ap
Sequence 1310, Appl
Sequence 131, Appl
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  Sequence 74, Application US/10927262A

Publication No. US20050137137A1

GENERAL INFORMATION:
APPLICANT: LANE, DAVID P
APPLICANT: BOTTGER, VOLKER
APPLICANT: BOTTGER, ANGELIKA
APPLICANT: BOTTGER, ANGELIKA
APPLICANT: PICKSLEY, STEVEN M.
APPLICANT: HOCHKEPPEL, HEINZ-KURT
APPLICANT: GARCIA-ECHEVERRIA, CARLOS
APPLICANT: CHENE, PATRICK
                                                                                                                                                                                                                  RESULT 2
US-10-927-262A-74
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US-09-214-371-74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PA
SEQ ID NO 74
LENGTH: 18
TYPE: PRT
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT.
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
                                                                                                                                                                                                                                                                                                   1 PPLSQETFSDLWKLLPEN 18
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Result No.

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28
29	65	65	65	65	65	66	66	66	66	66	66	66	66	66	66	66	66
61.9	61.9	61.9	61.9	61.9	61.9	62.9	62.9	62.9	62.9	62.9	62.9	62.9	62.9	62.9	62.9	62.9	62.9
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US-10-862-195-2239	US-10-793-943-3	US-10-215-982-55	US-09-952-680A-55	US-09-950-692-6	US-09-258-981-3	US-10-927-262A-39	US-10-340-179-18	US-10-339-712-38	US-10-927-262A-24	US-10-927-262A-17	US-10-645-784-143	US-10-645-784-131	US-10-653-048-143	US-10-653-048-131	US-10-666-696-143	US-10-666-696-131	US-10-645-761-143
Sequence 2239, Ap	Sequence 3, Appli	Sequence 55, Appl	Sequence 55, Appl	Sequence 6, Appli	Sequence 3, Appli	•	-	Sequence 38, Appl	•	Sequence 17, Appl	Sequence 143, App	•	•	Sequence 131, App	Sequence 143, App	Sequence 131, App	Sequence 143, App

ALIGNMENTS

APPLICANT: Lane, David APPLICANT: Bottger, Volker APPLICANT: Bottger, Angelica APPLICANT: Bottger, Angelica APPLICANT: Picksley, Stephen APPLICANT: Chene, Patrick APPLICANT: Hochkeppel, Heinz-Kurt APPLICANT: Garcia-Echeverria, Carlos APPLICANT: Furet, Pascal TITLE OF INVENTION: Inhibitors of the Interaction of P53 FILE REFERENCE: 4-20937/A/PCT CURRENT APPLICATION NUMBER: US/09/214,371B CURRENT FILING DATE: 1999-03-26 CURRENT FILING DATE: 1999-03-26 ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Description of Artificial Sequence:peptide 1 PPLSQETFSDLWKLLPEN 18 SEQ ID NOS: 83 PatentIn Ver. 2.0 Application US/09214371B Conservative 94.3%; Score 99; 100.0%; Pred. No. PCT/EP97/03549 , , Mismatches DB 3; 1 0 Length 18; Indels P53 and

MDM2

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Gaps

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RESULT 4
US-10-754-457-9
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Sequence 7, Application US/09732384
Patent No. US20020132977A1
PATENT NO US20020132977A1
GENERAL INFORMATION:
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Sequence 9, Application US/10754457
Publication No. US20040161798A1
GENERAL INFORMATION:
APPLICANT: KODADEK, THOMAS
TITLE OF INVENTION: METHODS AND COMPOSITIONS COMPRISING CAPTURE AGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Loc
Matches
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PRIOR FILING DATE: 1996-07-05
PRIOR APPLICATION NUMBER: GB 9707041.1
PRIOR FILING DATE: 1997-04-07
NUMBER OF SEQ ID NOS: 85
SOFTWARE: Patentin Ver. 3.2
SEQ ID NO 74
                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: INHIBITORS OF THE INTERACTION BETWEEN P53 AND MDM2
FILE REFERENCE: 39749.0002 APC CON
CURRENT APPLICATION NUMBER: US/10/927,262A
CURRENT FILING DATE: 2004-08-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/214,371 PRIOR FILING DATE: 1999-03-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Peptide
                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence:Protein OTHER INFORMATION: fragment not in inhibitory p53 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94.3%; Score 99; DB ilarity 100.0%; Pred. No. 9. Conservative 0; Mismatches
                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                              87.6%; Score 92; DB 3; I
100.0%; Pred. No. 1.2e-06;
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US-10-155-059-13
Sequence 13, Application US/10155059
Publication No. US20020147173A1
GENERAL INFORMATION:
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OTHER INFORMATION:
US-10-754-457-9
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                                                                                                                                RESULT 6
                                                                                                                                                                                                                                                                                                                                              US-10-927-262A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: LANE DAVID P
APPLICANT: BOTTGER, VOLKER
APPLICANT: BOTTGER, NIGELIKA
APPLICANT: BOTTGER, NIGELIKA
APPLICANT: PICKSLEY, STEVEN M.
APPLICANT: GACCIA-ECHEVERRIA,
APPLICANT: CHENE, PATRICK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/754,457
CURRENT FILING DATE: 2004-01-09
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1997-04-07
NUMBER OF SEQ ID NOS: 85
SOFTWARE: PatentIn Ver. 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/214,371
PRIOR FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
PRIOR APPLICATION NUMBER: GB 9614197.3
PRIOR APPLICATION NUMBER: GB 9707041.1
PRIOR APPLICATION NUMBER: GB 9707041.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: CHENE, PATRICK
APPLICANT: FURET, PASCAL
TITLE OF INVENTION: INHIBITORS OF THE INTERACTION BETWEEN P53 AND MDM2
FILE REFERENCE: 39749,0002 APC CON
CURRENT APPLICATION NUMBER: US/10/927,262A
CURRENT FILING DATE: 2004-08-25
CURRENT FILING DATE: 2004-08-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: UTSD:935US
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 19
Jost, Christine TITLE OF INVENTION: METHODS OF TREATMENT USING
                                   APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                   17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17;
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HOCHKEPPEL, HEINZ-KURT
GARCIA-ECHEVERRIA, CARLOS
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                                                                                                                                                                                                                                                                                   87.6%; Score 92; 100.0%; Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                   Mismatches
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. 1.2e-06;
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                                                                                                                                                                                                                                                                                                     Length 19;
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NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS:

NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE ANTIBODIES

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APPLICANT: Lane, Da
APPLICANT: Bottger,
APPLICANT: Bottger,
APPLICANT: Picksley,
APPLICANT: Chene, F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; SEQUENCE DESCRIPTION: SEQ ID NO: 13: US-10-155-059-13
SOFTWARE: P
SEQ ID NO 1
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Appli
Patent No. US2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                            APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REFERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
                                                                         PRIOR APPLICATION NUMBER: PCT PRIOR FILING DATE: 1997-07-04
                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

OPERATING SYSTEM: Windows

SOFTWARE: FastSEQ for Windows Version 2.0b

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/155,059

FILING DATE: 24-May-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             h 85.7%;
Similarity 88.9%;
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COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                           Application US/09214371B
JS20010018511A1
                                         PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 19 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Nixon Peabody LLP STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                           Bottger, Volker
Bottger, Angelica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Boston
                                                                                                                                                                                                                                                        Picksley, Stephen
Chene, Patrick
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                                                                                                                                                                                                                                                                                                                                      David
                                                                                                PCT/EP97/03549
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Pred. No. 2.3e-06
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GENERAL INFORMATION:

APPLICANT: Hupp, Theodore
APPLICANT: Dornan, David
TITLE OF INVENTION: Screening Method and Agents
FILE REFERENCE: 9013.54
CURRENT APPLICATION NUMBER: US/10/467,758
CURRENT FILING DATE: 2003-08-13
PRIOR APPLICATION NUMBER: PCT/GB02/00640
PRIOR APPLICATION NUMBER: GB 0103508.8
PRIOR FILING DATE: 2002-02-13
PRIOR FILING DATE: 2001-02-13
PRIOR FILING DATE: 2001-02-13
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US-10-211-088-141
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                                                                    PRIOR APPLICATION NUMBER: 60/341,589
PRIOR FILING DATE: 2001-12-13
NUMBER OF SEQ ID NOS: 366
SOFTWARE: PatentIn version 3.1
SEQ ID NO 141
LENGTH: 15
                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL GARY R.
APPLICANT: Premkunar, D. David
APPLICANT: Chen, Yih-Tai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn version 3.1
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                    Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 141,
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Best Local Similarity
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                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/211,088
CURRENT EILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/309,395
PRIOR FILING DATE: 2001-08-01
                                                                                                                                                                                                                                                                  TITLE OF INVENTION: No. US20030104479A1el Fusion Proteins And Assays For Molecular B FILE REFERENCE: 01-1022-US
OTHER INFORMATION: Binding domain
                                    TYPE: PRT ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 16
TYPE: PRT
ORGANISM: Homo sapiens
                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence:peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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Pred. No. 3.3e-06;
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; FEATURE:
; OTHER INFORMATION: peptide; amino acid residues 12-26 of human p53 protein US-11-019-894A-1
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Best Local S
Matches 15
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 4, Application US/10425970 Publication No. US20040052794A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.0 SEQ ID NO 4
                                                                                                                SOFTWARE: PatentIn version 3.2 SEQ ID NO 1
                                                                                                                                                                                                                                                                                        APPLICANT: Pincus, Matthew R.

TITLE OF INVENTION: PEPTIDES SELECTIVELY LETHAL TO THE MALIGNANT AND TRANSFORMED TITLE OF INVENTION: WAMMALIAN CELLS

FILE REFERENCE: 1181-17 CIP A
CURRENT APPLICATION NUMBER: US/11/019,894A
CURRENT FILING DATE: 2004-12-21
PRIOR APPLICATION NUMBER: 10/386,737
PRIOR ETLING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
PRIOR FILING DATE: 2003-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/425,970
CURRENT FILING DATE: 2003-04-30
PRIOR APPLICATION NUMBER: 09/581,472
PRIOR FILING DATE: 2000-08-14
PRIOR APPLICATION NUMBER: PCT/EP98/08560
PRIOR FILING DATE: 1998-12-14
PRIOR FILING DATE: 1998-12-14
                                                                                                                                                       PRIOR FILING DATE: 2002-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: EP 97403051.2
PRIOR FILING DATE: 1997-12-15
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: FRADE, Raymond TITLE OF INVENTION: p53 REGULATORY PROTEIN CALLED RB18A AND USES THEREOF FILE REFERENCE: P06781US00/BAS
                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/195,102 PRIOR FILING DATE: 2000-04-05
                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/363,785
                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 09/827,683
PRIOR FILING DATE: 2001-04-05
                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 15
TYPE: PRT
ORGANISM: Artificial
                                                                                            LENGTH:
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Pred. No.
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5. 4.1e-05;
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APPLICANT: Yuan, Zhi-Min

APPLICANT: Gu, JiJie

APPLICANT: Gu, JiJie

APPLICANT: Gu, JiJie

APPLICANT: Gu, JiJie

TITLE OF INVENTION: Inhibition of p53 Degradation

FILE REFERENCE: 21508-044

CURRENT APPLICATION NUMBER: US/09/732,384

CURRENT FILING DATE: 2000-12-07

PRIOR APPLICATION NUMBER: 60/169,816

PRIOR FILING DATE: 1999-12-08

NUMBER OF SEQ ID NOS: 10

SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: ; OTHER INFORMATION: oligomer US-11-128-722-14
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US-09-732-384-6
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                                                                                                                                                                                                          ; SEQ ID NO 6
; LENGTH: 15
; TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 6, Application US/09732384 Patent No. US20020132977A1
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Best Local
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FILE REFERENCE: 5793/1281
CURRENT APPLICATION NUMBER: US/11/128,722
CURRENT FILING DATE: 2005-05-13
PRIOR APPLICATION NUMBER: 60/574,964
PRIOR FILING DATE: 2004-05-27
NUMBER OF SEC
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APPLICANT: Chapman, Ross
TITLE OF INVENTION: METHODS FOR PREPARING INTERNALLY CONSTRAINED PEPTIDES
TITLE OF INVENTION: AND PEPTIDOMIMETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 34
                                                                                                                                 FEATURE: OTHER INFORMATION: Description of Artificial Sequence:Protein OTHER INFORMATION: fragment not in inhibitory p53 polypeptide
                                                                                                                                                                                        ORGANISM: Artificial Sequence
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                                                                     Local Similarity
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5 QETFSDLWKLLPEN 18
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                                                     73.3%; Score 77; DB 3; 100.0%; Pred. No. 0.0001 ative 0; Mismatches
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QETFSDLWKLLPEN 14

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APPLICANT: Celis, Estaban

APPLICANT: Celis, Estaban

APPLICANT: Keogh, Elissa

ITITLE OF INVENTION: Inducing Cellular Immune Responses To

ITITLE OF INVENTION: p53 Using Peptide And Nucleic Acid Compositions

ITITLE OF INVENTION: p53 Using Peptide And Nucleic Acid Compositions

CURRENT APPLICATION NUMBER: US/11/051,411

CURRENT APPLICATION NUMBER: US/09/458,297

PRIOR APPLICATION NUMBER: US/09/458,297

PRIOR APPLICATION NUMBER: US 09/017,735

PRIOR APPLICATION NUMBER: US 09/017,735

PRIOR APPLICATION NUMBER: PCT/US99/13789

PRIOR FILING DATE: 1998-02-03

PRIOR FILING DATE: 1998-06-17

PRIOR FILING DATE: 1998-06-17

PRIOR FILING DATE: 1998-06-17
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PRIOR PILLING DATE: 1999-12-10
PRIOR APPLICATION NUMBER: US 09/017,735
PRIOR FILLING DATE: 1998-02-03
PRIOR FILLING DATE: 1999-06-17
PRIOR FILLING DATE: 1999-06-17
PRIOR APPLICATION NUMBER: US 09/098,584
PRIOR FILLING DATE: 1998-06-17
NUMBER OF SEQ ID NOS: 1492
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 1109
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; OTHER INFORMATION: Synthetic Peptide
US-11-051-411-1109
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Best Local S
Matches 14
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APPLICANT:
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APPLICANT: Keogh, Elissa
TITLE OF INVENTION: Inducing Cellular Immune Responses To
TITLE OF INVENTION: p53 Using Peptide And Nucleic Acid Compositions
FILE REFERENCE: 2060.0120000
FILE REFERENCE: 1060.0120000
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APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
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CURRENT FILING DATE: 2005-02-07
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ORGANISM: Artificial Sequence
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Southwood, Scott
Chesnut, Robert
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Chesnut, Robert
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                                                                                                         ; OTHER INFORMATION: Synthetic Peptide US-11-051-411-1469
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SEQ ID NO 1469
LENGTH: 15
TYPE: PRT
ORGANISM: Artificial Sequence
                                                     Matches
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Best Local S
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2 PPLSQETFSDLWKL 15
                1 PPLSQETFSDLWKL 14
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Search completed: July 5, 2006, 23:04:06 Job time: 182 secs

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Result
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Listing first 45 summaries
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      July 5, 2006, 23:01:11 ; Search time 21 Seconds (without alignments) 24.277 Million cell updates/sec
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Copyright (c) 1993 - 2006
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US-10-953-613C-790
US-10-953-613C-792
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US-10-538-066-624
US-10-538-066-624
US-10-538-066-627
US-10-538-066-353
US-10-538-066-353
US-10-538-066-638
US-10-538-066-638
US-10-538-066-638
US-10-953-613C-893
US-10-953-613C-893
US-11-118-524-5
US-10-953-613C-894
US-10-953-613C-895
US-11-196-917A-85
US-11-953-613C-897
US-10-953-613C-897
US-10-953-613C-897
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Sequence 790,
Sequence 780,
Sequence 792,
Sequence 779,
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778, App
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781, App
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784, App
785, App
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787, App
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		ALIGNMENTS				
2112, Ap	Sequence 21		10	22.	24	45
789, App			10	22.	24	44
776, App		9- 7 US-11-134-871-776	19	23.8	25	4 .
292, App	Sequence 29	7 US-11-234	18	23.	25	42
3598, Ap		7 US-11-134-871-	18	23.	25	41
		7 US-11-134-	18	23.	25	40
	æ	6 US-10-953-613C-	17	23.	25	39
588, App		6 US-10-953-613C-5	12	23.	25	38
-		7 US-11-134-871-	11	23.	25	37
561, App		7 US-11-23	19	24.	26	36
		6 US-10-	12	24.	26	35
597, App		6 US-10-95	12	24.	26	34
		6 US-10-953-613C-	12	24.	26	33
	Sequence 58	6 US-10-953-613C-	12	24.	26	32
595, App		6 US-10-	11	24.	26	31
, Appli	Sequence 4,		11	24.	26	30
7, Appl	Sequence 77	7 US	10	24.	26	29
9, Appl	Sequence 29,		10	24.	26	28
907, App			17	25.	27	27
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CURRENT FILING DATE: 2004-09-29
PRIOR APPLICATION NUMBER: 60/507,231
PRIOR FILING DATE: 2003-09-30
NUMBER OF SEQ ID NOS: 1021
SOFTWARE: Patentin Ver 3.0
SEQ ID NO 778
LENGTH: 12
TYPE: PRT
ORGANISM: Homo sapiens
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SEQ ID NO 790
LENGTH: 12
TYPE: PRT
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Publication No. US20060127404A1
GENERAL INFORMATION:
APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon;
APPLICANT: Bernard; Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CEN5038 NP
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Matches
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                                                           CURRENT APPLICATION NUMBER: US/10/953,613C
CURRENT FILING DATE: 2004-09-29
PRIOR APPLICATION NUMBER: 60/507,231
PRIOR FILING DATE: 2003-09-30
NUMBER OF SEQ ID NOS: 1021
SOFTWARE: Patentin Ver 3.0
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CURRENT APPLICATION NUMBER: US/10/953,613C
CURRENT FILING DATE: 2004-09-29
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TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
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100.0%; Pred. No. 0.
cive 0; Mismatches
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                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:

APPLICANT: Huang, Chichi; Heavner, George; Kniq
APPLICANT: Bernard; Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES,
FILE REFERENCE: CEN5038 NP
CURRENT APPLICATION UNMERER: US/10/953,613C
CURRENT FILLING DATE: 2004-09-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Publication No. US20060127404A1 GENERAL INFORMATION:
                                                                                                                                                                                                     SOFTWARE: PatentIn Ver 3.0 SEQ ID NO 792
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Best Local
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TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
                                                                                                                                                                                                                                       CURRENT FILING DATE: 2004-09-29
PRIOR APPLICATION NUMBER: 60/507,231
PRIOR FILING DATE: 2003-09-30
NUMBER OF SEQ ID NOS: 1021
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ORGANISM: Homo sapiens
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rd;Nesspor; Thomas
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                                                                Score 61; DB 6; Pred. No. 0.003; 0; Mismatches
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Pred. No.
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RESULT 5

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; Sequence 624, Application US/10538066
; Publication No. US20060094649A1
; GENERAL INFORMATION:
; APPLICANT: Epimmune Inc.
; TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7,
; TITLE OF INVENTION: Peptides and Compositions
; FILE REFERENCE: 2060.015PC06
; CURRENT APPLICATION NUMBER: US/10/538,066
; CURRENT FILING DATE: 2005-06-09
; PRIOR APPLICATION NUMBER: US 60/432,017
; PRIOR FILING DATE: 2002-11-10
; NUMBER OF SEQ ID NOS: 767
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US-10-538-066-624
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SOFTWARE: PatentIn Ver 3.0
SEQ ID NO 791
LENGTH: 12
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SEQ ID NO 779
LENGTH: 12
TYPE: PRT
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon; APPLICANT: Bernard; Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CEM5038 NP
CURRENT APPLICATION NUMBER: US/10/953,613C
CURRENT FILING DATE: 2004-09-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon; APPLICANT: Bernard; Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CUN5038 NP
CURRENT APPLICATION NUMBER: US/10/953,613C
CURRENT FILING DATE: 2004-09-29
PRIOR APPLICATION NUMBER: 60/507,231
PRIOR FILING DATE: 2003-09-30
NUMBER OF SEQ ID NOS: 1021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/507,231 PRIOR FILING DATE: 2003-09-30
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ORGANISM: Homo sapiens
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Pred. No. 0.0042;
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Pred. No. 0.0042;
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NUMBER OF SEQ ID NOS: 767
SOFTWARE: Patentin version 3.2
SEQ ID NO 600
LENGTH: 10
TYPE: PRT
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-953-613C-781
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US-10-538-066-600
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US-10-538-066-624
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                                                                                                                                                                                                     Publication No. US20060127404A1

GENERAL INFORMATION:
APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon;
APPLICANT: Bernard, Nesspor; Thomas

TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CENS038 NP
CURRENT APPLICATION NUMBER: US/10/953,613C

CURRENT FILING DATE: 2004-09-29

PRIOR APPLICATION NUMBER: 60/507,231

PRIOR FILING DATE: 2003-09-30

PRIOR FILING DATE: 2003-09-30
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Best Local S
Matches 10
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SOFTWARE: PatentIn Ver 3.0
SEQ ID NO 781
                                               Matches
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Best Local Similarity
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Best Local :
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TITLE OF INVENTION: Paptides and Compositions
FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
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TYPE: PRT
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QETFSDLWKLLP 16
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Conservative
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; Pred. No. 0.019;
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Pred. No.
                                                          Score 55; DB 6;
Pred. No. 0.023;
                                               Mismatches
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APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon; ITILE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES; FILE REFERENCE: CENSO38 NP; CURRENT APPLICATION NUMBER: US/10/953,613C; CURRENT FILING DATE: 2004-09-29; PRIOR APPLICATION NUMBER: 60/507,231; PRIOR APPLICATION NUMBER: 60/507,231; PRIOR FILING DATE: 2003-09-30; NUMBER of SEQ ID NOS: 1021; SOFTWARE: Patentin Ver 3.0; SEQ ID NO 793
                                                                                                                                                RESULT 12
US-10-538-066-623
                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens US-10-538-066-353
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US-10-538-066-353
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Sequence 623, Application US/10538066
Publication No. US20060094649A1
GENERAL INFORMATION:
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7,
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Publication No. US20060094649A1
GENERAL INFORMATION:
APPLICANT: Epimmune.Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 767
SOFTWARE: PatentIn version 3.2
SEQ ID NO 353
LENGTH: 9
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TYPE: PRT
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o. US20060094649A1
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100.0%; Pr
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k; Pred. No. 1e+
0; Mismatches
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Pred. No. 0.023;
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1e+05;
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                                                            and -B44 Tumor Associated Antigen
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CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
SOFTWARE: Patentin version 3.2
SEQ ID NO 623
LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
US-10-538-066-623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, TITLE OF INVENTION: Peptides and Compositions FILE REFERENCE: 2060.015PC06
CURRENT APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US 60/432,017
PRIOR FILING DATE: 2002-12-10
NUMBER OF SEQ ID NOS: 767
SOFTWARE: PatentIn version 3.2
SEQ ID NO 354
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Best Local Similarity
Matches 8; Conserva
                                                   ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-538-066-622
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                                                                                              ; Sequence 622, Application US/10538066
; Publication No. US20060094649A1
GENERAL INFORMATION:
APPLICANT: Epimmune Inc.
TITLE OF INVENTION: HLA-A1, -A2, -A3, -A24, -B7, and -B44 Tumor Associated Antigen
TITLE OF INVENTION: Peptides and Compositions
FILE REFERENCE: 2060.015PC06
FULE REFERENCE: 2060.015PC06
CURRENT FILING DATE: 2005-06-09
PRIOR APPLICATION NUMBER: US/10/538,066
CURRENT FILING DATE: 2002-12-10
NUMBER OF 5EQ ID NOS: 767
SOFTWARE: PatentIn version 3.2
SEQ ID NO 622
LENGTH: 9
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Query Match
Best Local Similarity
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Best Local (
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TYPE: PRT
ORGANISM: Homo sapiens
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Local Similarity 87.5%;
res 7; Conservative
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35.2%;
100.0%;
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Pred. No. 1e+05;
Score 37;
Pred. No.
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DB 6;
1e+05;
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US-10-953-613C-777
Search completed: July 5, 2006, 23:04:32 Job time : 22 secs
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                                                                                                                                                                                                             ; ORGANISM: Homo sapiens US-10-953-613C-777
                                                                                                                                                                                                                                                                                                                Sequence 777, Application US/10953613C
Publication No. US20060127404A1

GENERAL INFORMATION:
APPLICANT: Huang, Chichi; Heavner, George; Knight, David; Ghrayeb, John; Scallon APPLICANT: Bernard; Nesspor; Thomas
TITLE OF INVENTION: HINGE CORE MIMETIBODIES, COMPOSITIONS, METHODS AND USES
FILE REFERENCE: CEMS038 NP
CURRENT APPLICATION NUMBER: US/10/953,613C
CURRENT FILING DATE: 2004-09-29
PRIOR APPLICATION NUMBER: 60/507,231
PRIOR FILING DATE: 2003-99-30
NUMBER OF SEQ ID NOS: 1021
                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver 3.0
SEQ ID NO 777
LENGTH: 6
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Best Local Similarity
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100.0%;
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b; Pred. No. 1e+
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GenCore version 5.1.9
Copyright (c) 1993 - 2006 Biocceleration Ltd.

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Result
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Searched:
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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     protein search, using sw model
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length: 19
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Match Length
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1: /EMC_Celerra_SIDS3/ptodata/2/iaa/5_COMB.pep:*

2: /EMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*

3: /EMC_Celerra_SIDS3/ptodata/2/iaa/7_COMB.pep:*

4: /EMC_Celerra_SIDS3/ptodata/2/iaa/H_COMB.pep:*

5: /EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*

6: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*

7: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
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(without alignments)
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US-09-681-472B-4
US-09-581-472B-4
US-08-277-660A-1
US-08-277-660A-1
US-08-424-957-1
US-08-424-957-20
US-09-035-686-20
US-09-035-686-20
US-09-035-686-21
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US-09-695-437A-8
US-09-695-437A-42
US-09-701-080C-27
US-09-695-437A-20
US-09-695-437A-64
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Sequence 7, Appli
Sequence 13, Appli
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Sequence 40, Appli
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Sequence 41, Appli
Sequence 64, Appli
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ALIGNMENTS	US-09-258-981-3 US-09-511-2048-3 US-09-515-692-6 US-09-695-437A-11 PCT-US95-02856-6 US-09-695-437A-12 US-09-695-437A-3 US-09-428-0828-133 US-09-428-0828-145 US-09-428-0828-144 US-09-428-0828-144 US-09-428-0828-132 US-09-428-0828-132 US-09-695-437A-3 US-09-695-637-660A-9 US-08-277-660A-9 US-09-035-686-17 US-09-603-052-5
	Sequence 3, Appli Sequence 6, Appli Sequence 6, Appli Sequence 11, Appli Sequence 16, Appli Sequence 16, Appli Sequence 21, Appl Sequence 23, Appl Sequence 13, Appl Sequence 13, Appl Sequence 13, Appl Sequence 13, Appl Sequence 14, Appl Sequence 22, Appl Sequence 33, Appl Sequence 34, Appl Sequence 17, Appl Sequence 37, Appl Sequence 27, Appl Sequence 5, Appl Sequence 17, Appl Sequence 17, Appl Sequence 5, Appli Sequence 5, Appli

RESULT 1

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GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
ITITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence:Protein ; OTHER INFORMATION: fragment not in inhibitory p53 polypeptide US-09-732-384-7
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US-09-081-975-13
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                                                                                                                     Patent No. 6451979

GENERAL INFORMATION:
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MES-1, ANTIBODIES AND PROTE
TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                     Best Loc
Matches
                                                                                                                                                                                                                                 Sequence 13, Application US/09081975 Patent No. 6451979
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 19
ADUAL STREET: 10 CITY: Boston
                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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17; Conserv
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                                                101 Federal Street
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/09581472B
Patent No. 6818744
GENERAL INFORMATION:
APPLICANT: FRADE, Raymond
TITLE OF INVENTION: p53 REGULATORY PROTEIN CALLED RB18A AND USES THEREOF
FILE REFERENCE: P06781US00/BAS
CURRENT APPLICATION NUMBER: US/09/581,472B
CURRENT FLING DATE: 2000-08-14
PRIOR APPLICATION NUMBER: PCT/EP98/08560
PRIOR FILING DATE: 1998-12-14
                                                                                                                                                                                                                             SEQ ID NO 4
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Best Local S
                                                               Matches
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Best Local (
                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: EP 97403051.2
PRIOR FILING DATE: 1997-12-15
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn version 3.0
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SEQUENCE CHARACTERISTICS
                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                              FEATURE:
                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 47-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/0.
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORWATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino a STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM CON OPERATING SYSTEM: SOFTWARE: FastSE(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0: FILING DATE: 12-MAY-1998 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 02110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                            l Similarity
15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 PPLSQETFSDLWKLLPEN 18
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                             PLSQETFSDLWKLLP 16
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PLSQETFSDLWKLLP 15
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                                                               Conservative
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IBM Compatible
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                                                                             77.1%; Score 81; 100.0%; Pred. No.
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BER: 47400
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                                                               Mismatches
                                                        DB 2; Lu...
3. 2.3e-06;
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                                                               Indels
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RESULT 4

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US-08-277-660A-1
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Best Local Similarity 100.0%; F
Marches 14; Conservative 0;
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Patent No. 5702908
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                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENT: Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660;
FILING DATE: 20-UUL-1994
CLASSIFICATION: 514
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                ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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LENGTH: 15 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                       NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach,
STREET: Four Embarcadero Ce
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                                                                                                                                                                                                                                                                    APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding
TITLE OF INVENTION: Protein and Therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
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CITY: San Francisco
                                                                                                                CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
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0; Mismatches
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Center, Suite 3400
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Center, Suite 3400
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Application
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RESULT 6
US-08-424-957-1
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GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
TITLE OF INVENTION: 50
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Best Local (
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                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELEPHONE: (415) 781-1999
TELEPHONE: (415) 388-3249
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TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                    TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
                                                  SEQUENCE CHARACTERISTICS
LENGTH: 15 amino acid
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NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-0
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hes 14; Conservative
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TOPOLOGY:
                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: United States
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                                 amino acid
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unknown
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100.0%; Prr
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Pred. No.
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o. le-05;
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; Sequence 1, Application US/09035686
; Patent No. 6153391
; GENERAL INFORMATION:
; APPLICANT: Picksley, Steven M.
; APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of TITLE OF INVENTION: Protein and TITLE OF SEQUENCES: 50
                                                                                                                                                        RESULT 8
US-09-035-686-1
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PATENT NO. 5770377
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 398-324
TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Dreger, Walter H. REGISTRATION NUMBER: 24 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: San Francisco
STATE: California
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     Interruption of Binding of MDM2 and P53
Protein and Therapeutic Application Thereof
50
                                                                                                                                                                                                                                                                                                         73.3%; Score 77; DB 1; 100.0%; Pred. No. 1e-05; tive 0; Mismatches
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CORRESPONDENCE ADDRESS:

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Best Local S
Matches 14
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Patent No. 6153391
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
                                                      COMPUTER READABLE FORM:
MEDICIN TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPAX: (415) 398-3249
TELEX: 410 22222
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                            APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                 STREET: Four Embarca
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: Four Embarca
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Dreger, Walter H. REGISTRATION NUMBER: 24
                     FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                          TRY: United States 94111-4187
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                                                                                                                                                                                                                                                                                           E: Flehr, Hohbach, Test, Albritton & Herbert Four Embarcadero Center, Suite 3400
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                                                                                                       #1.30
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GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, Jülie
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION UNMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 6
INVERTAL 15
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US-08-277-660A-5
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US-09-732-384-6
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                                                                                                                     Sequence 5, Application US/08277660A Patent No. 5702908
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 6, Application US/09732384
Patent No. 6831155
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              GENERAL INFORMATION:
APPLICANT: PicksLey, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding
TITLE OF INVENTION: Protein and Therapeutic
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (415) 781-198
TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08)
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,11
                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 15
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LENGTH: 15 amino acids
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100.0%; |
les 14; Conservative 0;
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                                                                                                                                                                                                                                                                                                                        Local
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                                                                                                                                                                                                                                                                                                       14;
                                                                                                                                                                                                                              5 QETFSDLWKLLPEN 18
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1 QETFSDLWKLLPEN 14
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                                                                                                                                                                                                                                                                                                                        73.3%; Score 77; DB 2; 100.0%; Pred. No. 1e-05;
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Pred. No.
                                                                                                                                                                                                                                                                                                         Mismatches
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                                 of MDM2 and Application
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ADDRESSEE:

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US-08-424-957-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE DOCKET NUMBER: A-60:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELERAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. 5770377
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21,
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Best Local :
                                                                                                                                              ZIP: 94111-11.

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTION Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

CURRENT APPLICATION UNMBER: US/08/424,957
ATTORNEY AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-6
                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Flahr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                CITY: San Francisco
STATE: California
                                                                                                                                      FILING DATE: 1
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 20-JU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: United States ZIP: 94111-4187
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14; Conserv
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    A-61228/WHD
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Pred. No. 3e-05;
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RESULT 13
US-09-035-686-21
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Matches
                Query Match
Best Local Similarity
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                                                                                                                                                                       TELEFAX: (415) 398-324
TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                        FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 781-
TELEFAX: (415) 398-32-
TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/424,957
FILLING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
APPLICATION NUMBER: US 08/277,660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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STATE: California
Thired Stat
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TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Picksley, Steven M. APPLICANT: Lane, David P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: United States ZIP: 94111-4187
                                                                                                     STRANDEDNESS:
                                                                                                                         TYPE:
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Similarity 93.3%;
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Conservative
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              Score 74; DB 2;
Pred. No. 3e-05;
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Pred. No. 3e-05;
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Mismatches
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Sequence 63, Application US/09695437A
Patent No. 6803203
GENERAL INFORMATION:
APPLICANT: Brookhaven Science Associates
APPLICANT: Anderson, Carl W
APPLICANT: Connelly, Margery A
TITLE OF INVENTION: DNA-PK Assay
FILE REFERENCE: BSA 01-02
CURRENT FILING DATE: 2000-10-24
PRIOR APPLICATION NUMBER: US/09/695,437A
PRIOR APPLICATION NUMBER: US/09/695,437A
PRIOR APPLICATION NUMBER: US/09/695,437A
PRIOR APPLICATION NUMBER: US/09/695,437A
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US-09-081-975-23
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US-09-695-437A-63
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Best Local :
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APPLICANT: Kaelin, William
APPLICANT: JOSE, Christine
APPLICANT: JOSE, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
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LENGTH: 16 amino acids
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CITY: Boston
STATE: MA
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TYPE: amino acid
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ZIP: 02110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: 1i
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linear
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Search completed: July Job time : 51 secs
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                                                                                                                                                            , OTHER INFORMATION: DNA-PK synthetic substrate based on human US-09-695-437A-63
                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 08/132,284
PRIOR FILING DATE: 1993-10-06
NUMBER OF SEQ ID NOS: 64
SOFTWARE: PatentIn version 3.1
SEQ ID NO 63
LENGTH: 18
TYPE: PRT
                                                                                                                      Best
                                                                                                        Matches
                                                                                                                                   Query Match
                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE:
                                                                                                       Local Similarity 93.3 hes 14; Conservative
                                                                    2 PLSQETFSDLWKLLP 16
                                                  PLSQETFSGLWKLLP
            5, 2006, 23:00:59
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                                                                                                                      Score 74;
Pred. No.
                                                                                                           Mismatches
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3.7e-05;
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